



#6

SEQUENCE LISTING

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 CCTCTCTGTC aagCTCTTC CTGTGGGCTT cctgaggCCc ccaAGAAATGG aatGGTGTtT 6780
 ggcAaggAGT acacAGTGGG aaccaAGGCC atgtACAGCT gcAGTGAAGG ctaccACCTC 6840
 caggcaggCGC ctgaggCCAC tgCAGAGTGT ctggACACAG gcCTATGGAG caACCGCAAT 6900
 gtcCcaccAC agtGTgtCCG tgagtCCTCG ggCAATGGAG gcGGGtCTGT gacttGtCCT 6960
 gatgtcAGTA gcatcAGCgt ggAGCATGGC cgATGGAGGC ttATCTTGA gACACAGTAT 7020
 cagttCCAGG cccAGCTGAT gctcatCTGT gaccCTGGCT actactatac tggccAAAGG 7080
 gtcatCCGCT gtcaggCCAA tggCAAATGG AGCCTCGGGG actCTACGCC cacCTGCCGA 7140
 atcatCTCTCt gtggAGAGCT cccgattCCC cccaATGGCC accGcatCgg aacACTGTCT 7200
 gtctacGGGG caacAGCCAT ctTCTCCTGC aattCCGGAT acACACTGtT gggCTCCAGG 7260
 gtgcgtgAGT gcatGGCCAA tgggCTCTGG agtGGCTCTG aagtCCGCTG CCTTGCCACT 7320
 cagaccaAGC tccactCCAT ttTCTATAAG ctccTCTTCG atgtACTCTC ttccccatCC 7380
 ctcaccaaAG ctggACACTG tgggACTCCT gagCCCAATTG tcaACGGACA catCAATGGG 7440
 gagaactaca gctacGGGG cagtGTGGT taccaATGCA atgCTGGCTT ccgcCTGATC 7500
 ggcAtgtCTG tgCgcAtCTG ccAGCAGGAT catCActGGT cgggCAAGAC ccCTTCTGT 7560
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 atggCTGAGG gggCTGCTAG gtCCCAATGc ctggCCAGCG ggCAATGGAG tgACATGCTG 7800
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<210> 4
 <211> 2669
 <212> PRT
 <213> Homo sapiens

<400> 4
 Met Ala Gly Ala Pro Pro Pro Ala Leu Leu Leu Pro Cys Ser Leu Ile
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Ser Glu Leu Val Lys Lys Gln Ile Glu Leu Lys Ser Arg Gly Val Lys
 35 40 45

Leu Met Pro Ser Lys Asp Asn Ser Gln Lys Thr Ser Val Leu Thr Gln
 50 55 60

Val Gly Val Ser Gln Gly His Asn Met Cys Pro Asp Pro Gly Ile Pro
 65 70 75 80

Glu Arg Gly Lys Arg Leu Gly Ser Asp Phe Arg Leu Gly Ser Ser Val

85	90	95
Gln Phe Thr Cys Asn Glu Gly Tyr Asp Leu Gln	Gly Ser Lys Arg Ile	
100	105	110
Thr Cys Met Lys Val Ser Asp Met Phe Ala Ala Trp	Ser Asp His Arg	
115	120	125
Pro Val Cys Arg Ala Arg Met Cys Asp Ala His Leu	Arg Gly Pro Ser	
130	135	140
Gly Ile Ile Thr Ser Pro Asn Phe Pro Ile Gln	Tyr Asp Asn Asn Ala	
145	150	155
His Cys Val Trp Ile Ile Thr Ala Leu Asn Pro Ser	Lys Val Ile Lys	
165	170	175
Leu Ala Phe Glu Glu Phe Asp Leu Glu Arg Gly Tyr	Asp Thr Leu Thr	
180	185	190
Val Gly Asp Gly Gly Gln Asp Gly Asp Gln Lys	Thr Val Leu Tyr Met	
195	200	205
Ser Gln Asn Ala Cys Ser Asp Ser Pro His Thr Pro	Gly Ser Arg Ile	
210	215	220
Pro Glu Ser Met Ser Gly Asp Ile Trp Arg Gln	Lys Trp Thr Val Leu	
225	230	235
240		
Glu Ile Cys Arg Asp Ile Ser Ser Asp Ala Arg Ser	Gly Ser Val	
245	250	255
Arg Lys Ser Pro Lys Thr Ser Asn Ala Val Glu	Leu Val Ala Pro Gly	
260	265	270
Thr Glu Ile Glu Gln Gly Ser Cys Gly Asp Pro	Gly Ile Pro Ala Tyr	
275	280	285
Gly Arg Arg Glu Gly Ser Arg Phe His His Gly	Asp Thr Leu Lys Phe	
290	295	300
Glu Cys Gln Pro Ala Phe Glu Leu Val Gly Gln	Lys Ala Ile Thr Cys	
305	310	315
320		
Gln Lys Asn Asn Gln Trp Ser Ala Lys Lys Pro	Gly Cys Val Phe Ser	
325	330	335
Cys Phe Phe Asn Phe Thr Ser Pro Ser Gly Val	Val Val Leu Ser Pro Asn	
340	345	350
Tyr Pro Glu Asp Tyr Gly Asn His Leu His Cys	Val Trp Leu Ile Leu	
355	360	365
Ala Arg Pro Glu Ser Arg Ile His Leu Ala Phe	Asn Asp Ile Asp Val	
370	375	380
Glu Pro Gln Phe Asp Phe Leu Val Ile Lys Asp	Gly Ala Thr Ala Glu	

385	390	395	400
Ala Pro Val Leu Gly Thr Phe Ser Gly Asn Gln Leu Pro Ser Ser Ile			
405		410	415
Thr Ser Ser Gly His Val Ala Arg Leu Glu Phe Gln Thr Asp His Ser			
420		425	430
Thr Gly Lys Arg Gly Phe Asn Ile Thr Phe Thr Thr Phe Arg His Asn			
435		440	445
Glu Cys Pro Asp Pro Gly Val Pro Val Asn Gly Lys Arg Phe Gly Asp			
450		455	460
Ser Leu Gln Leu Gly Ser Ser Ile Ser Phe Leu Cys Asp Glu Gly Phe			
465		470	475
Leu Gly Thr Gln Gly Ser Glu Thr Ile Thr Cys Val Leu Lys Glu Gly			
485		490	495
Ser Val Val Trp Asn Ser Ala Val Leu Arg Cys Glu Ala Pro Cys Gly			
500		505	510
Gly His Leu Thr Ser Pro Ser Gly Thr Ile Leu Ser Pro Gly Trp Pro			
515		520	525
Gly Phe Tyr Lys Asp Ala Leu Ser Cys Ala Trp Val Ile Glu Ala Gln			
530		535	540
Pro Gly Tyr Pro Ile Lys Ile Thr Phe Asp Arg Phe Lys Thr Glu Val			
545		550	555
Asn Tyr Asp Thr Leu Glu Val Arg Asp Gly Arg Thr Tyr Ser Ala Pro			
565		570	575
Leu Ile Gly Val Tyr His Gly Thr Gln Val Pro Gln Phe Leu Ile Ser			
580		585	590
Thr Ser Asn Tyr Leu Tyr Leu Leu Phe Ser Thr Asp Lys Ser His Ser			
595		600	605
Asp Ile Gly Phe Gln Leu Arg Tyr Glu Thr Ile Thr Leu Gln Ser Asp			
610		615	620
His Cys Leu Asp Pro Gly Ile Pro Val Asn Gly Gln Arg His Gly Asn			
625		630	635
Asp Phe Tyr Val Gly Ala Leu Val Thr Phe Ser Cys Asp Ser Gly Tyr			
645		650	655
Thr Leu Ser Asp Gly Glu Pro Leu Glu Cys Glu Pro Asn Phe Gln Trp			
660		665	670
Ser Arg Ala Leu Pro Ser Cys Glu Ala Leu Cys Gly Gly Phe Ile Gln			
675		680	685
Gly Ser Ser Gly Thr Ile Leu Ser Pro Gly Phe Pro Asp Phe Tyr Pro			

690	695	700
Asn Asn Leu Asn Cys Thr Trp Ile Ile Glu Thr Ser His Gly Lys Gly		
705	710	715
Val Phe Phe Thr Phe His Thr Phe His Leu Glu Ser Gly His Asp Tyr		
725	730	735
Leu Leu Ile Thr Glu Asn Gly Ser Phe Thr Gln Pro Leu Arg Gln Leu		
740	745	750
Thr Gly Ser Arg Leu Pro Ala Pro Ile Ser Ala Gly Leu Tyr Gly Asn		
755	760	765
Phe Thr Ala Gln Val Arg Phe Ile Ser Asp Phe Ser Met Ser Tyr Glu		
770	775	780
Gly Phe Asn Ile Thr Phe Ser Glu Tyr Asp Leu Glu Pro Cys Glu Glu		
785	790	795
Pro Glu Val Pro Ala Tyr Ser Ile Arg Lys Gly Leu Gln Phe Gly Val		
805	810	815
Gly Asp Thr Leu Thr Phe Ser Cys Phe Pro Gly Tyr Arg Leu Glu Gly		
820	825	830
Thr Ala Arg Ile Thr Cys Leu Gly Gly Arg Arg Arg Leu Trp Ser Ser		
835	840	845
Pro Leu Pro Arg Cys Val Ala Glu Cys Gly Asn Ser Val Thr Gly Thr		
850	855	860
Gln Gly Thr Leu Leu Ser Pro Asn Phe Pro Val Asn Tyr Asn Asn Asn		
865	870	875
His Glu Cys Ile Tyr Ser Ile Gln Thr Gln Pro Gly Lys Gly Ile Gln		
885	890	895
Leu Lys Ala Arg Ala Phe Glu Leu Ser Glu Gly Asp Val Leu Lys Val		
900	905	910
Tyr Asp Gly Asn Asn Ser Ala Arg Leu Leu Gly Val Phe Ser His		
915	920	925
Ser Glu Met Met Gly Val Thr Leu Asn Ser Thr Ser Ser Ser Leu Trp		
930	935	940
Leu Asp Phe Ile Thr Asp Ala Glu Asn Thr Ser Lys Gly Phe Glu Leu		
945	950	955
His Phe Ser Ser Phe Glu Leu Ile Lys Cys Glu Asp Pro Gly Thr Pro		
965	970	975
Lys Phe Gly Tyr Lys Val His Asp Glu Gly His Phe Ala Gly Ser Ser		
980	985	990
Val Ser Phe Ser Cys Asp Pro Gly Tyr Ser Leu Arg Gly Ser Glu Glu		

995	1000	1005
Leu Leu Cys Leu Ser Gly Glu Arg Arg Thr Trp Asp Arg Pro Leu Pro		
1010	1015	1020
Thr Cys Val Ala Glu Cys Gly Gly Thr Val Arg Gly Glu Val Ser Gly		
1025	1030	1035
Gln Val Leu Ser Pro Gly Tyr Pro Ala Pro Tyr Glu His Asn Leu Asn		
1045	1050	1055
Cys Ile Trp Thr Ile Glu Ala Glu Ala Gly Cys Thr Ile Gly Leu His		
1060	1065	1070
Phe Leu Val Phe Asp Thr Glu Glu Val His Asp Val Leu Arg Ile Trp		
1075	1080	1085
Asp Gly Pro Val Glu Ser Gly Val Leu Leu Lys Glu Leu Ser Gly Pro		
1090	1095	1100
Ala Leu Pro Lys Asp Leu His Ser Thr Phe Asn Ser Val Val Leu Gln		
1105	1110	1115
Phe Ser Thr Asp Phe Phe Thr Ser Lys Gln Gly Phe Ala Ile Gln Phe		
1125	1130	1135
Ser Val Ser Thr Ala Thr Ser Cys Asn Asp Pro Gly Ile Pro Gln Asn		
1140	1145	1150
Gly Ser Arg Ser Gly Asp Ser Trp Glu Ala Gly Asp Ser Thr Val Phe		
1155	1160	1165
Gln Cys Asp Pro Gly Tyr Ala Leu Gln Gly Ser Ala Glu Ile Ser Cys		
1170	1175	1180
Val Lys Ile Glu Asn Arg Phe Phe Trp Gln Pro Ser Pro Pro Thr Cys		
1185	1190	1195
Ile Ala Pro Cys Gly Gly Asp Leu Thr Gly Pro Ser Gly Val Ile Leu		
1205	1210	1215
Ser Pro Asn Tyr Pro Glu Pro Tyr Pro Pro Gly Lys Glu Cys Asp Trp		
1220	1225	1230
Lys Val Thr Val Ser Pro Asp Tyr Val Ile Ala Leu Val Phe Asn Ile		
1235	1240	1245
Phe Asn Leu Glu Pro Gly Tyr Asp Phe Leu His Ile Tyr Asp Gly Arg		
1250	1255	1260
Asp Ser Leu Ser Pro Leu Ile Gly Ser Phe Tyr Gly Ser Gln Leu Pro		
1265	1270	1275
Gly Arg Ile Glu Ser Ser Ser Asn Ser Leu Phe Leu Ala Phe Arg Ser		
1285	1290	1295
Asp Ala Ser Val Ser Asn Ala Gly Phe Val Ile Asp Tyr Thr Glu Asn		

1300	1305	1310
Pro Arg Glu Ser Cys Phe Asp Pro Gly Ser Ile Lys Asn Gly Thr Arg		
1315	1320	1325
Val Gly Ser Asp Leu Lys Leu Gly Ser Ser Val Thr Tyr Tyr Cys His		
1330	1335	1340
Gly Gly Tyr Glu Val Glu Gly Thr Ser Thr Leu Ser Cys Ile Leu Gly		
1345	1350	1355
Pro Asp Gly Lys Pro Val Trp Asn Asn Pro Arg Pro Val Cys Thr Ala		
1365	1370	1375
Pro Cys Gly Gly Gln Tyr Val Gly Ser Asp Gly Val Val Leu Ser Pro		
1380	1385	1390
Asn Tyr Pro Gln Asn Tyr Thr Ser Gly Gln Ile Cys Leu Tyr Phe Val		
1395	1400	1405
Thr Val Pro Lys Asp Tyr Val Val Phe Gly Gln Phe Ala Phe Phe His		
1410	1415	1420
Thr Ala Leu Asn Asp Val Val Glu Val His Asp Gly His Ser Gln His		
1425	1430	1435
Ser Arg Leu Leu Ser Ser Leu Ser Gly Ser His Thr Gly Glu Ser Leu		
1445	1450	1455
Pro Leu Ala Thr Ser Asn Gln Val Leu Ile Lys Phe Ser Ala Lys Gly		
1460	1465	1470
Leu Ala Pro Ala Arg Gly Phe His Phe Val Tyr Gln Ala Val Pro Arg		
1475	1480	1485
Thr Ser Ala Thr Gln Cys Ser Ser Val Pro Glu Pro Arg Tyr Gly Lys		
1490	1495	1500
Arg Leu Gly Ser Asp Phe Ser Val Gly Ala Ile Val Arg Phe Glu Cys		
1505	1510	1515
Asn Ser Gly Tyr Ala Leu Gln Gly Ser Pro Glu Ile Glu Cys Leu Pro		
1525	1530	1535
Val Pro Gly Ala Leu Ala Gln Trp Asn Val Ser Ala Pro Thr Cys Val		
1540	1545	1550
Val Pro Cys Gly Gly Asn Leu Thr Glu Arg Arg Gly Thr Ile Leu Ser		
1555	1560	1565
Pro Gly Phe Pro Glu Pro Tyr Leu Asn Ser Leu Asn Cys Val Trp Lys		
1570	1575	1580
Ile Val Val Pro Glu Gly Ala Gly Ile Gln Ile Gln Val Val Ser Phe		
1585	1590	1595
Val Thr Glu Gln Asn Trp Asp Ser Leu Glu Val Phe Asp Gly Ala Asp		

1605	1610	1615
Asn Thr Val Thr Met Leu Gly Ser Phe Ser Gly Thr Thr Val Pro Ala		
1620	1625	1630
Leu Leu Asn Ser Thr Ser Asn Gln Leu Tyr Leu His Phe Tyr Ser Asp		
1635	1640	1645
Ile Ser Val Ser Ala Ala Gly Phe His Leu Glu Tyr Lys Thr Val Gly		
1650	1655	1660
Leu Ser Ser Cys Pro Glu Pro Ala Val Pro Ser Asn Gly Val Lys Thr		
1665	1670	1675
Gly Glu Arg Tyr Leu Val Asn Asp Val Val Ser Phe Gln Cys Glu Pro		
1685	1690	1695
Gly Tyr Ala Leu Gln Gly His Ala His Ile Ser Cys Met Pro Gly Thr		
1700	1705	1710
Val Arg Arg Trp Asn Tyr Pro Pro Pro Leu Cys Ile Ala Gln Cys Gly		
1715	1720	1725
Gly Thr Val Glu Glu Met Glu Gly Val Ile Leu Ser Pro Gly Phe Pro		
1730	1735	1740
Gly Asn Tyr Pro Ser Asn Met Asp Cys Ser Trp Lys Ile Ala Leu Pro		
1745	1750	1755
1760		
Val Gly Phe Gly Ala His Ile Gln Phe Leu Asn Phe Ser Thr Glu Pro		
1765	1770	1775
Asn His Asp Tyr Ile Glu Ile Arg Asn Gly Pro Tyr Glu Thr Ser Arg		
1780	1785	1790
Met Met Gly Arg Phe Ser Gly Ser Glu Leu Pro Ser Ser Leu Leu Ser		
1795	1800	1805
Thr Ser His Glu Thr Thr Val Tyr Phe His Ser Asp His Ser Gln Asn		
1810	1815	1820
Arg Pro Gly Phe Lys Leu Glu Tyr Gln Ala Tyr Glu Leu Gln Glu Cys		
1825	1830	1835
1840		
Pro Asp Pro Glu Pro Phe Ala Asn Gly Ile Val Arg Gly Ala Gly Tyr		
1845	1850	1855
Asn Val Gly Gln Ser Val Thr Phe Glu Cys Leu Pro Gly Tyr Gln Leu		
1860	1865	1870
Thr Gly His Pro Val Leu Thr Cys Gln His Gly Thr Asn Arg Asn Trp		
1875	1880	1885
Asp His Pro Leu Pro Lys Cys Glu Val Pro Cys Gly Gly Asn Ile Thr		
1890	1895	1900
Ser Ser Asn Gly Thr Val Tyr Ser Pro Gly Phe Pro Ser Pro Tyr Ser		

1905	1910	1915	1920
Ser Ser Gln Asp Cys Val Trp Leu Ile Thr Val Pro Ile Gly His Gly			
1925		1930	1935
Val Arg Leu Asn Leu Ser Leu Leu Gln Thr Glu Pro Ser Gly Asp Phe			
1940		1945	1950
Ile Thr Ile Trp Asp Gly Pro Gln Gln Thr Ala Pro Arg Leu Gly Val			
1955		1960	1965
Phe Thr Arg Ser Met Ala Lys Lys Thr Val Gln Ser Ser Ser Asn Gln			
1970		1975	1980
Val Leu Leu Lys Phe His Arg Asp Ala Ala Thr Gly Gly Ile Phe Ala			
1985	1990	1995	2000
Ile Ala Phe Ser Ala Tyr Pro Leu Thr Lys Cys Pro Pro Pro Thr Ile			
2005		2010	2015
Leu Pro Asn Ala Glu Val Val Thr Glu Asn Glu Glu Phe Asn Ile Gly			
2020		2025	2030
Asp Ile Val Arg Tyr Arg Cys Leu Pro Gly Phe Thr Leu Val Gly Asn			
2035		2040	2045
Glu Ile Leu Thr Cys Lys Leu Gly Thr Tyr Leu Gln Phe Glu Gly Pro			
2050		2055	2060
Pro Pro Ile Cys Glu Val His Cys Pro Thr Asn Glu Leu Leu Thr Asp			
2065	2070	2075	2080
Ser Thr Gly Val Ile Leu Ser Gln Ser Tyr Pro Gly Ser Tyr Pro Gln			
2085		2090	2095
Phe Gln Thr Cys Ser Trp Leu Val Arg Val Glu Pro Asp Tyr Asn Ile			
2100		2105	2110
Ser Leu Thr Val Glu Tyr Phe Leu Ser Glu Lys Gln Tyr Asp Glu Phe			
2115		2120	2125
Glu Ile Phe Asp Gly Pro Ser Gly Gln Ser Pro Leu Leu Lys Ala Leu			
2130		2135	2140
Ser Gly Asn Tyr Ser Ala Pro Leu Ile Val Thr Ser Ser Ser Asn Ser			
2145		2150	2155
Val Tyr Leu Arg Trp Ser Ser Asp His Ala Tyr Asn Arg Lys Gly Phe			
2165		2170	2175
Lys Ile Arg Tyr Ser Ala Pro Tyr Cys Ser Leu Pro Arg Ala Pro Leu			
2180		2185	2190
His Gly Phe Ile Leu Gly Gln Thr Ser Thr Gln Pro Gly Gly Ser Ile			
2195		2200	2205
His Phe Gly Cys Asn Ala Gly Tyr Arg Leu Val Gly His Ser Met Ala			

2210	2215	2220
Ile Cys Thr Arg His Pro Gln Gly Tyr His Leu Trp Ser Glu Ala Ile		
2225	2230	2235
Pro Leu Cys Gln Ala Leu Ser Cys Gly Leu Pro Glu Ala Pro Lys Asn		
2245	2250	2255
Gly Met Val Phe Gly Lys Glu Tyr Thr Val Gly Thr Lys Ala Met Tyr		
2260	2265	2270
Ser Cys Ser Glu Gly Tyr His Leu Gln Ala Gly Ala Glu Ala Thr Ala		
2275	2280	2285
Glu Cys Leu Asp Thr Gly Leu Trp Ser Asn Arg Asn Val Pro Pro Gln		
2290	2295	2300
Cys Val Arg Glu Ser Ser Gly Asn Gly Gly Ser Val Thr Cys Pro		
2305	2310	2315
Asp Val Ser Ser Ile Ser Val Glu His Gly Arg Trp Arg Leu Ile Phe		
2325	2330	2335
Glu Thr Gln Tyr Gln Phe Gln Ala Gln Leu Met Leu Ile Cys Asp Pro		
2340	2345	2350
Gly Tyr Tyr Tyr Thr Gly Gln Arg Val Ile Arg Cys Gln Ala Asn Gly		
2355	2360	2365
Lys Trp Ser Leu Gly Asp Ser Thr Pro Thr Cys Arg Ile Ile Ser Cys		
2370	2375	2380
Gly Glu Leu Pro Ile Pro Pro Asn Gly His Arg Ile Gly Thr Leu Ser		
2385	2390	2395
Val Tyr Gly Ala Thr Ala Ile Phe Ser Cys Asn Ser Gly Tyr Thr Leu		
2405	2410	2415
Val Gly Ser Arg Val Arg Glu Cys Met Ala Asn Gly Leu Trp Ser Gly		
2420	2425	2430
Ser Glu Val Arg Cys Leu Ala Thr Gln Thr Lys Leu His Ser Ile Phe		
2435	2440	2445
Tyr Lys Leu Leu Phe Asp Val Leu Ser Ser Pro Ser Leu Thr Lys Ala		
2450	2455	2460
Gly His Cys Gly Thr Pro Glu Pro Ile Val Asn Gly His Ile Asn Gly		
2465	2470	2475
Glu Asn Tyr Ser Tyr Arg Gly Ser Val Val Tyr Gln Cys Asn Ala Gly		
2485	2490	2495
Phe Arg Leu Ile Gly Met Ser Val Arg Ile Cys Gln Gln Asp His His		
2500	2505	2510
Trp Ser Gly Lys Thr Pro Phe Cys Val His Val Lys Gln Gln Leu Leu		

2515

2520

2525

Leu Leu Leu Leu Leu Cys Asp Asp Asp Asp Asp Glu Asp Asp Gly
2530 2535 2540

Ser Gly Ala Ile Thr Cys Gly His Pro Gly Asn Pro Val Asn Gly Leu
2545 2550 2555 2560

Thr Gln Gly Asn Gln Phe Asn Leu Asn Asp Val Val Lys Phe Val Cys
2565 2570 2575

Asn Pro Gly Tyr Met Ala Glu Gly Ala Ala Arg Ser Gln Cys Leu Ala
2580 2585 2590

Ser Gly Gln Trp Ser Asp Met Leu Pro Thr Cys Arg Ile Ile Asn Cys
2595 2600 2605

Thr Asp Pro Gly His Gln Glu Asn Ser Val Arg Gln Val His Ala Ser
2610 2615 2620

Gly Pro His Arg Phe Ser Phe Gly Thr Thr Val Ser Tyr Arg Cys Asn
2625 2630 2635 2640

His Gly Phe Tyr Leu Leu Gly Thr Pro Val Leu Ser Cys Gln Gly Asp
2645 2650 2655

Gly Thr Trp Asp Arg Pro Arg Pro Gln Cys Leu Cys Lys
2660 2665

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<211> 1464

<212> DNA

<213> Homo sapiens

<400> 5

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Glu Ile Arg Ala Asp Ala His Val Arg Gly Tyr Val Gly Glu Lys Ile
35 40 45

Lys Leu Lys Cys Thr Phe Lys Ser Thr Ser Asp,Val Thr Asp Lys Leu
50 55 60

Thr Ile Asp Trp Thr Tyr Arg Pro Pro Ser Ser Ser His Thr Val Ser
65 70 75 80

Ile Phe His Tyr Gln Ser Phe Gln Tyr Pro Thr Thr Ala Gly Thr Phe
85 90 95

Arg Asp Arg Ile Ser Trp Val Gly Asn Val Tyr Lys Gly Asp Ala Ser
100 105 110

Ile Ser Ile Ser Asn Pro Thr Ile Lys Asp Asn Gly Thr Phe Ser Cys
115 120 125

Ala Val Lys Asn Pro Pro Asp Val His His Asn Ile Pro Met Thr Glu
130 135 140

Leu Thr Val Thr Glu Arg Gly Phe Gly Thr Met Leu Ser Ser Val Ala
145 150 155 160

Leu Leu Ser Ile Leu Val Phe Val Pro Ser Ala Val Val Val Ala Leu
165 170 175

Leu Leu Val Arg Met Gly Arg Lys Ala Ala Gly Leu Lys Lys Arg Ser
180 185 190

Arg Ser Gly Tyr Lys Lys Ser Ser Ile Glu Val Ser Asp Asp Thr Asp
195 200 205

Gln Glu Glu Glu Glu Ala Cys Met Ala Arg Leu Val Ser Val Ala Leu
210 215 220

Ser Ala Trp Ile Gln Thr Met Lys Arg His Ile Asp Glu Ser Leu Tyr
225 230 235 240

Asp Thr Arg Arg Val Thr
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<212> DNA
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Pro Glu Gly Ser Arg Trp Val Pro Pro Asp Ser Ala Cys Ser Ser Cys

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Ser Cys Ala Gln Pro Arg Gln Gly Pro His Asp Cys Cys Pro Gln Cys		
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Ser Asp Cys Glu His Glu Gly Arg Lys Tyr Glu Pro Gly Glu Ser Phe		
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Gln Pro Gly Ala Asp Pro Cys Glu Val Cys Ile Cys Glu Pro Gln Pro		
115	120	125
Glu Gly Pro Pro Ser Leu Arg Cys His Arg Arg Gln Cys Pro Ser Leu		
130	135	140
Val Gly Cys Pro Pro Ser Gln Leu Leu Pro Pro Gly Pro Gln His Cys		
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Cys Pro Thr Cys Ala Glu Ala Leu Ser Asn Cys Ser Glu Gly Leu Leu		
165	170	175
Gly Ser Glu Leu Ala Pro Pro Asp Pro Cys Tyr Thr Cys Gln Cys Gln		
180	185	190
Asp Leu Thr Trp Leu Cys Ile His Gln Ala Cys Pro Glu Leu Ser Cys		
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Pro Leu Ser Glu Arg His Thr Pro Pro Gly Ser Cys Cys Pro Val Cys		
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Arg Glu Cys Val Val Glu Ala Glu Gly Arg Arg Val Ala Asp Gly Glu		
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Ser Trp Arg Asp Pro Ser Asn Ala Cys Ile Ala Cys Thr Cys His Arg		
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Gly His Val Glu Cys His Leu Glu Glu Cys Gln Ala Leu Ser Cys Pro		
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Tyr Val Glu Leu Arg Gly His Thr Val Ile Leu His Ala Gln Pro Gly		
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Arg Asp Pro His Gly Leu Gln Gly Pro Gly Val Gly Pro Ala Pro Gly		
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Lys Ala Pro Ala Gly Leu Ser Ala Ala Gln Gln Val Thr Ala Leu Gln		

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Arg Pro Cys Pro Arg Ala Pro Cys Ala His Pro Leu Pro Gly Thr Cys		
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Cys Pro Asn Asp Cys Ser Gly Cys Ala Phe Gly Gly Lys Glu Tyr Pro		
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Asp Gly Ser Val Ser Cys Gln Arg Leu Pro Cys Pro Pro Ala Pro Cys			
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Ser Gly Asn Glu Ser Gln His Arg Leu Asp Phe Gln Leu Met Leu Lys
 50 55 60

Ile Arg Asp Thr Leu Tyr Ile Ala Gly Arg Asp Gln Val Tyr Thr Val
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Thr Trp Arg Ser Arg Gln Gln Asp Arg Glu Asn Cys Ala Met Lys Gly
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Lys His Lys Asp Glu Cys His Asn Phe Ile Lys Val Phe Val Pro Arg
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 145 150 155 160
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 165 170 175
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 370 375 380
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 385 390 395 400
 Ser Ala Val Pro Pro Ile Ala Asp Glu Pro Trp Phe Thr Lys Thr Arg
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Tyr Gln Asn Tyr Thr Val Ile Phe Val Gly Ser Glu Ala Gly Met Val
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1 5 10 15

Gln Leu Arg Ala Val Ser Phe Pro Glu Asp Asp Glu Pro Leu Asn Thr
 20 25 30

Val Asp Tyr His Cys Lys Ser Ser Arg Gln Tyr Pro Val Phe Arg Gly
35 40 45

Arg Pro Ser Gly Asn Glu Ser Gln His Arg Leu Asp Phe Gln Leu Met
50 55 60

Leu Lys Ile Arg Asp Thr Leu Tyr Ile Ala Gly Arg Asp Gln Val Tyr
65 70 75 80

Thr Val Asn Leu Asn Glu Met Pro Lys Thr Glu Val Ile Trp Gln Gln
85 90 95

Lys Leu Thr Trp Arg Ser Arg Gln Gln Asp Arg Glu Asn Cys Ala Met
100 105 110

Lys Gly Lys His Lys Asp Glu Cys His Asn Phe Ile Lys Val Phe Val
115 120 125

Pro Arg Asn Asp Glu Met Val Phe Val Cys Gly Thr Asn Ala Phe Asn
130 135 140

Pro Met Cys Arg Tyr Tyr Arg Val Ser Thr Leu Glu Tyr Asp Gly Glu
145 150 155 160

Glu Ile Ser Gly Leu Ala Arg Cys Pro Phe Asp Ala Arg Gln Thr Asn
165 170 175

Val Ala Leu Phe Ala Asp Gly Lys Leu Tyr Ser Ala Thr Val Ala Asp
180 185 190

Phe Leu Ala Ser Asp Ala Val Ile Tyr Arg Ser Met Gly Asp Gly Ser
 195 200 205
 Ala Leu Arg Thr Ile Lys Tyr Asp Ser Lys Trp Ile Lys Glu Pro His
 210 215 220
 Phe Leu His Ala Ile Glu Tyr Gly Asn Tyr Val Tyr Phe Phe Arg
 225 230 235 240
 Glu Ile Ala Val Glu His Asn Asn Leu Gly Lys Ala Val Tyr Ser Arg
 245 250 255
 Val Ala Arg Ile Cys Lys Asn Asp Met Gly Gly Ser Gln Arg Val Leu
 260 265 270
 Glu Lys His Trp Thr Ser Phe Leu Lys Ala Arg Leu Asn Cys Ser Val
 275 280 285
 Pro Gly Asp Ser Phe Phe Tyr Phe Asp Val Leu Gln Ser Ile Thr Asp
 290 295 300
 Ile Ile Gln Ile Asn Gly Ile Pro Thr Val Val Gly Val Phe Thr Thr
 305 310 315 320
 Gln Leu Asn Ser Ile Pro Gly Ser Ala Val Cys Ala Phe Ser Met Asp
 325 330 335
 Asp Ile Glu Lys Val Phe Lys Gly Arg Phe Lys Glu Gln Lys Thr Pro
 340 345 350
 Asp Ser Val Trp Thr Ala Val Pro Glu Asp Lys Val Pro Lys Pro Arg
 355 360 365
 Pro Gly Cys Cys Ala Lys His Gly Leu Ala Glu Ala Tyr Lys Thr Ser
 370 375 380
 Ile Asp Phe Pro Asp Glu Thr Leu Ser Phe Ile Lys Ser His Pro Leu
 385 390 395 400
 Met Asp Ser Ala Val Pro Pro Ile Ala Asp Glu Pro Trp Phe Thr Lys
 405 410 415
 Thr Arg Val Arg Tyr Arg Leu Thr Ala Ile Ser Val Asp His Ser Ala
 420 425 430
 Gly Pro Tyr Gln Asn Tyr Thr Val Ile Phe Val Gly Ser Glu Ala Gly
 435 440 445
 Met Val Leu Lys Val Leu Ala Lys Thr Ser Pro Phe Ser Leu Asn Asp
 450 455 460
 Ser Val Leu Leu Glu Glu Ile Glu Ala Tyr Asn His Ala Lys Cys Ser
 465 470 475 480
 Ala Glu Asn Glu Glu Asp Lys Lys Val Ile Ser Leu Gln Leu Asp Lys
 485 490 495

Asp His His Ala Leu Tyr Val Ala Phe Ser Ser Cys Ile Ile Arg Ile
 500 505 510
 Pro Leu Ser Arg Cys Glu Arg Tyr Gly Ser Cys Lys Lys Ser Cys Ile
 515 520 525
 Ala Ser Arg Asp Pro Tyr Cys Gly Trp Leu Ser Gln Gly Ser Cys Gly
 530 535 540
 Arg Val Thr Pro Asn His Ser Ala Glu Gly Tyr Glu Gln Asp Thr Glu
 545 550 555 560
 Phe Gly Asn Thr Ala His Leu Gly Asp Cys His Ala Tyr Glu Pro Tyr
 565 570 575
 Glu Gly Arg Val Gly Ser Leu Lys Ala Ile Cys Tyr Leu Leu Phe
 580 585 590
 Leu Lys Ser Thr Leu Phe Thr Leu Ser His Val Ser Ile Ser Gly Val
 595 600 605
 Arg Trp Glu Val Gln Ser Gly Glu Ser Asn Gln Met Val His Met Asn
 610 615 620
 Val Leu Ile Thr Cys Val Phe Ala Ala Phe Val Leu Gly Ala Phe Ile
 625 630 635 640
 Ala Gly Val Ala Val Tyr Cys Tyr Arg Asp Met Phe Val Arg Lys Asn
 645 650 655
 Arg Lys Ile His Lys Asp Ala Glu Ser Ala Gln Ser Cys Thr Asp Ser
 660 665 670
 Ser Gly Ser Phe Ala Lys Leu Asn Gly Leu Phe Asp Ser Pro Val Lys
 675 680 685
 Glu Tyr Gln Gln Asn Ile Asp Ser Pro Lys Leu Tyr Ser Asn Leu Leu
 690 695 700
 Thr Ser Arg Lys Glu Leu Pro Pro Asn Gly Asp Thr Lys Ser Met Val
 705 710 715 720
 Met Asp His Arg Gly Gln Pro Pro Glu Leu Ala Ala Leu Pro Thr Pro
 725 730 735
 Glu Ser Thr Pro Val Leu His Gln Lys Thr Leu Gln Ala Met Lys Ser
 740 745 750
 His Ser Glu Lys Ala His Gly His Gly Ala Ser Arg Lys Glu Thr Pro
 755 760 765
 Gln Phe Phe Pro Ser Ser Pro Pro Pro His Ser Pro Leu Ser His Gly
 770 775 780
 His Ile Pro Ser Ala Ile Val Leu Pro Asn Ala Thr His Asp Tyr Asn
 785 790 795 800

Thr Ser Phe Ser Asn Ser Asn Ala His Lys Ala Glu Lys Lys Leu Gln
 805 810 815
 Asn Ile Asp His Pro Leu Thr Lys Ser Ser Ser Lys Arg Asp His Arg
 820 825 830
 Arg Ser Val Asp Ser Arg Asn Thr Leu Asn Asp Leu Leu Lys His Leu
 835 840 845
 Asn Asp Pro Asn Ser Asn Pro Lys Ala Ile Met Gly Asp Ile Gln Met
 850 855 860
 Ala His Gln Asn Leu Met Leu Asp Pro Met Gly Ser Met Ser Glu Val
 865 870 875 880
 Pro Pro Lys Val Pro Asn Arg Glu Ala Ser Leu Tyr Ser Pro Pro Ser
 885 890 895
 Thr Leu Pro Arg Asn Ser Pro Thr Lys Arg Val Asp Val Pro Thr Thr
 900 905 910
 Pro Gly Val Pro Met Thr Ser Leu Glu Arg Gln Arg Gly Tyr His Lys
 915 920 925
 Asn Ser Ser Gln Arg His Ser Ile Ser Ala Met Pro Lys Asn Leu Asn
 930 935 940
 Ser Pro Asn Gly Val Leu Leu Ser Arg Gln Pro Ser Met Asn Arg Gly
 945 950 955 960
 Gly Tyr Met Pro Thr Pro Thr Gly Ala Lys Val Asp Tyr Ile Gln Gly
 965 970 975
 Thr Pro Val Ser Val His Leu Gln Pro Ser Leu Ser Arg Gln Ser Ser
 980 985 990
 Tyr Thr Ser Asn Gly Thr Leu Pro Arg Thr Gly Leu Lys Arg Thr Pro
 995 1000 1005
 Ser Leu Lys Pro Asp Val Pro Pro Lys Pro Ser Phe Val Pro Gln Thr
 1010 1015 1020
 Pro Ser Val Arg Pro Leu Asn Lys Tyr Thr Tyr
 1025 1030 1035

 <210> 13
 <211> 2191
 <212> DNA
 <213> Homo sapiens

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 gatgatgaac cccttaataac tgtcgactat cactgtaagt cgtctaggca atatccgggt 180
 tttagaggac gcccttcagg caatgaatcg cagcacagggc tggactttca gctgatgttg 240

aaaattcgag acacactta tattgctggc agggatcaag tttatacagt aaacttaaat 300
gaaatgccca aaacagaagt aatatggcaa cagaaactga catggcgatc aagacaacag 360
gatcgagaaa actgtgctat gaaaggcaag cataaaagatg aatgccacaa ctatcaaa 420
gtatgttgc caagaaacga tgagatggtt ttgtttgtg gtaccaatgc attcaatccc 480
atgtgttagat actacagggt aagtaccta gaatatgtat gggagaataat tagtggctg 540
gcaagatgcc catttgatgc cagacaaacc aatgtgccc tcttgcgtga tggaaagctg 600
tattctgcca cagtggctga cttctggcc agcgatgccc ttatattatcg aagcatgggt 660
gatggatctg cccttcgcac aataaaatat gattccaaat ggataaaaaga gccacacttt 720
cttcatgcca tagaatatgg aaactatgtc tatttcttct ttcgagaataat cgctgtcgaa 780
cataataatt taggcaaggc tggatattcc cgcgtggccc gcatatgtaa aaacgacatg 840
ggtggttccc agcgggtcct ggagaaacac tggacttcat ttctaaaggc tcggctgaac 900
tggatgtcc ctggagattc gttttctac ttgtatgttc tgcagtctat tacagacata 960
atacaatca atggcatccc cactgtggc ggggttta ccacgcagct caatagcatc 1020
cctgggtctg ctgtctgtc atttagcatg gatgacattt aaaaagtatt caaaggacgg 1080
tttaaggaac agaaaactcc agattctgtt tggacagcag ttcccgaaga caaagtgcac 1140
aagccaaggc ctggctgtt tgcaaaacac ggcctgccc aagcttataa aacccatc 1200
gatttcccgg atgaaactct gtcattcatc aaatctcatc ccctgtatgaa ctctgccc 1260
ccaccattt ccgatgagcc ctgggtcaca aagactcggg tcaggtacag actgacggcc 1320
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gaagctggca tggacttta agttctggc aagaccagtc ctttctctt gaacgacagc 1440
gtattactgg aagagattga agcctacaac catgcaaaatg gcagtgtca gaatgaggaa 1500
gacaaaaagg tcatctcatt acagttggat aaagatcacc acgcttataa tgtggcgttc 1560
tctagctgca ttatcccat cccccctcagt cgctgtgagc gttatggatc atgtaaaaag 1620
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cgatggaaag tccagtttgg agagtccaaac cagatggtcc acatgaatgt cctcatcacc 1920
tgtgtctttt ctgctttgt tttggggca ttcattgcag gtgtggcagt atactgctat 1980
cgagacatgt ttgttcggaa aaacagaaag atccataaaatg atgcagatgc cggccagtc 2040
tgcacagact ccagtttggaa ttttgcctaaa ctgaatggtc tctttgacag ccctgtcaag 2100
gaataccaac agaatatttgc ttctcctaaa ctgtatagta acctgctaaatc cagtcggaaa 2160
gagcacgaat tcagcggccg ctgaattctat g 2191

<210> 14
<211> 712
<212> PRT
<213> Homo sapiens

<400> 14
Met Arg Val Phe Leu Leu Cys Ala Tyr Ile Leu Leu Leu Met Val Ser
1 5 10 15

Gln Leu Arg Ala Val Ser Phe Pro Glu Asp Asp Glu Pro Leu Asn Thr
20 25 30

Val Asp Tyr His Cys Lys Ser Ser Arg Gln Tyr Pro Val Phe Arg Gly
35 40 45

Arg Pro Ser Gly Asn Glu Ser Gln His Arg Leu Asp Phe Gln Leu Met
50 55 60

Leu Lys Ile Arg Asp Thr Leu Tyr Ile Ala Gly Arg Asp Gln Val Tyr
65 70 75 80

Thr Val Asn Leu Asn Glu Met Pro Lys Thr Glu Val Ile Trp Gln Gln

85	90	95
Lys Leu Thr Trp Arg Ser Arg Gln Gln Asp Arg Glu Asn Cys Ala Met		
100	105	110
Lys Gly Lys His Lys Asp Glu Cys His Asn Phe Ile Lys Val Phe Val		
115	120	125
Pro Arg Asn Asp Glu Met Val Phe Val Cys Gly Thr Asn Ala Phe Asn		
130	135	140
Pro Met Cys Arg Tyr Tyr Arg Val Ser Thr Leu Glu Tyr Asp Gly Glu		
145	150	155
160		
Glu Ile Ser Gly Leu Ala Arg Cys Pro Phe Asp Ala Arg Gln Thr Asn		
165	170	175
Val Ala Leu Phe Ala Asp Gly Lys Leu Tyr Ser Ala Thr Val Ala Asp		
180	185	190
Phe Leu Ala Ser Asp Ala Val Ile Tyr Arg Ser Met Gly Asp Gly Ser		
195	200	205
Ala Leu Arg Thr Ile Lys Tyr Asp Ser Lys Trp Ile Lys Glu Pro His		
210	215	220
Phe Leu His Ala Ile Glu Tyr Gly Asn Tyr Val Tyr Phe Phe Arg		
225	230	235
240		
Glu Ile Ala Val Glu His Asn Asn Leu Gly Lys Ala Val Tyr Ser Arg		
245	250	255
Val Ala Arg Ile Cys Lys Asn Asp Met Gly Gly Ser Gln Arg Val Leu		
260	265	270
Glu Lys His Trp Thr Ser Phe Leu Lys Ala Arg Leu Asn Cys Ser Val		
275	280	285
Pro Gly Asp Ser Phe Phe Tyr Phe Asp Val Leu Gln Ser Ile Thr Asp		
290	295	300
Ile Ile Gln Ile Asn Gly Ile Pro Thr Val Val Gly Val Phe Thr Thr		
305	310	315
320		
Gln Leu Asn Ser Ile Pro Gly Ser Ala Val Cys Ala Phe Ser Met Asp		
325	330	335
Asp Ile Glu Lys Val Phe Lys Gly Arg Phe Lys Glu Gln Lys Thr Pro		
340	345	350
Asp Ser Val Trp Thr Ala Val Pro Glu Asp Lys Val Pro Lys Pro Arg		
355	360	365
Pro Gly Cys Cys Ala Lys His Gly Leu Ala Glu Ala Tyr Lys Thr Ser		
370	375	380
Ile Asp Phe Pro Asp Glu Thr Leu Ser Phe Ile Lys Ser His Pro Leu		

385	390	395	400
Met Asp Ser Ala Val Pro Pro Ile Ala Asp Glu Pro Trp Phe Thr Lys			
405		410	415
Thr Arg Val Arg Tyr Arg Leu Thr Ala Ile Ser Val Asp His Ser Ala			
420		425	430
Gly Pro Tyr Gln Asn Tyr Thr Val Ile Phe Val Gly Ser Glu Ala Gly			
435		440	445
Met Val Leu Lys Val Leu Ala Lys Thr Ser Pro Phe Ser Leu Asn Asp			
450		455	460
Ser Val Leu Leu Glu Glu Ile Glu Ala Tyr Asn His Ala Lys Cys Ser			
465		470	475
Ala Glu Asn Glu Glu Asp Lys Lys Val Ile Ser Leu Gln Leu Asp Lys			
485		490	495
Asp His His Ala Leu Tyr Val Ala Phe Ser Ser Cys Ile Ile Arg Ile			
500		505	510
Pro Leu Ser Arg Cys Glu Arg Tyr Gly Ser Cys Lys Lys Ser Cys Ile			
515		520	525
Ala Ser Arg Asp Pro Tyr Cys Gly Trp Leu Ser Gln Gly Ser Cys Gly			
530		535	540
Arg Val Thr Pro Gly Met Leu Leu Leu Thr Glu Asp Phe Phe Ala Phe			
545		550	555
His Asn His Ser Ala Glu Gly Tyr Glu Gln Asp Thr Glu Phe Gly Asn			
565		570	575
Thr Ala His Leu Gly Asp Cys His Glu Ile Leu Pro Thr Ser Thr Thr			
580		585	590
Pro Asp Tyr Lys Ile Phe Gly Gly Pro Thr Ser Gly Val Arg Trp Glu			
595		600	605
Val Gln Ser Gly Glu Ser Asn Gln Met Val His Met Asn Val Leu Ile			
610		615	620
Thr Cys Val Phe Ala Ala Phe Val Leu Gly Ala Phe Ile Ala Gly Val			
625		630	635
Ala Val Tyr Cys Tyr Arg Asp Met Phe Val Arg Lys Asn Arg Lys Ile			
645		650	655
His Lys Asp Ala Glu Ser Ala Gln Ser Cys Thr Asp Ser Ser Gly Ser			
660		665	670
Phe Ala Lys Leu Asn Gly Leu Phe Asp Ser Pro Val Lys Glu Tyr Gln			
675		680	685
Gln Asn Ile Asp Ser Pro Lys Leu Tyr Ser Asn Leu Leu Thr Ser Arg			

690

695

700

Lys Glu His Glu Phe Ser Gly Arg
705 710

<210> 15
<211> 3196
<212> DNA
<213> *Homo sapiens*

<400> 15
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gatgatgaac cccttaatac tgctgactat cactgtaaat cgtcttaggca atatcccggtt 180
tttagaggac gcccttcagg caatgaatcg cagcacagggc tggactttca gctgtatggtt 240
aaaattcgag acacacttta tattgtctggc agggatcaag tttatacagt aaacttaaat 300
gaaatgccc aAACAGAAGT aatatggcaa cagaaactgta catggcgatc aagacaacag 360
gatcgagaaa actgtgctat gaaaggcaag cataaaagatg aatgccacaa ctttatcaaa 420
gtatTTGTTc caagaaacga tgagatggtt ttgtttgtt gtaccaatgc attcaatccc 480
atgtgttagat actacagggt aagtaccttta gaatatgtatg gggaaagaat tagtggcctg 540
gcaagatgcc catttgatgc cagacaaacc aatgttgc ctttgctga tgggaagctg 600
tattctgcca cagtggctga cttctggcc agcgatgccc ttatttatcg aagcatgggt 660
gatggatctg cccttcgcac aataaaatat gattccaaat ggataaaaga gcccacactt 720
cttcatgcca tagaatatgg aaactatgtc tatttcttct ttgagaaat cgctgtcgaa 780
cataataatt taggcaaggc tgtgtattcc cgcgtggccc gcataatgtaa aaacgacatg 840
ggtgttccc agcgggtcct ggagaaacac tggacttcat ttctaaaggc tcggctgaac 900
tggctgtcc ctggagattc gttttctac ttgtatggc tgcaatgtat tacagacata 960
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cctggttctg ctgtctgtgc atttagcatg gatgacattt aaaaagtatt caaaggacgg 1080
ttaagaac agaaaaactcc agattctgtt tggacagcag ttccgaaaga aacgtccatc 1140
aagccaaggc ctggctgtt tgcaaaacac ggccttgccc aagcttataa ctctggcattt 1200
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gaagctggca tggtaacttaa agttctggca aagaccagtc ctttctttaa gaacgacagc 1440
gtattactgg aagagattga agcctacaac catgcaaaat gcagtgctga gaatgaggaa 1500
gacaaaaagg tcatctcatt acagttggat aaagatcacc acgcattata tggcgttc 1560
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cgatggaaag tccagtcgtt agagtcac cagatggtcc acatgaatgt cctcatcacc 1920
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cgagacatgt ttgttcggaa aaacagaaag atccataaaat atgcagagtc cggccagtc 2040
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gaataccaac agaatattga ttctccat cttgtatagta acctgtcaac cagtcggaaa 2160
gagctaccac ccaatggaga ttctaaatcc atggtaatgg accatcgagg gcaacccctt 2220
gagttggctg ctcttcctac tcctgagttt acacccgtgc ttcaccagaa gaccctgcag 2280
gccatgaaga gccactcaga aaaggcccat ggcctatggag cttcaaggaa agaaaccctt 2340
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aatgacccaa atagtaaccc caaagccatc atgggagaca tccagatggc acaccagaac 2640
ttaatgtgg atccccatggg atcgatgtct gaggtccac cttaaatgtccc taaccgggag 2700

gcatcgctat	actcccctcc	ttcaactctc	cccagaaaata	gcccaaccaa	gcgagtggat	2760
gtccccacca	ctcctggagt	cccaatgact	tctctggaaa	gacaaggagg	ttatcacaaa	2820
aattcctccc	agaggcactc	tatatctgct	atgcctaaaa	acttaaactc	accaaatggt	2880
gttttgttat	ccagacagcc	tagtatgaac	cgtggaggat	atatggccac	ccccactggg	2940
gcgaagggtgg	actatattca	gggaacacca	gtgagtgttc	atctgcagcc	ttccctctcc	3000
agacagagca	gctacaccag	taatggcaact	cttccttagga	cgggactaaa	gaggacgccg	3060
tccttaaaac	ctgacgtgcc	accaaagcct	tcctttgttc	ctcaaacccc	atctgtcaga	3120
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tccgtgttgt	tgagag					3196

<210> 16
<211> 1032
<212> PRT
<213> Homo sapiens

<400> 16
 Met Arg Val Phe Leu Leu Cys Ala Tyr Ile Leu Leu Leu Met Val Ser
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 Gln Leu Arg Ala Val Ser Phe Pro Glu Asp Asp Glu Pro Leu Asn Thr
 20 25 30

 Val Asp Tyr His Cys Lys Ser Ser Arg Gln Tyr Pro Val Phe Arg Gly
 35 40 45

 Arg Pro Ser Gly Asn Glu Ser Gln His Arg Leu Asp Phe Gln Leu Met
 50 55 60

 Leu Lys Ile Arg Asp Thr Leu Tyr Ile Ala Gly Arg Asp Gln Val Tyr
 65 70 75 80

 Thr Val Asn Leu Asn Glu Met Pro Lys Thr Glu Val Ile Trp Gln Gln
 85 90 95

 Lys Leu Thr Trp Arg Ser Arg Gln Gln Asp Arg Glu Asn Cys Ala Met
 100 105 110

 Lys Gly Lys His Lys Asp Glu Cys His Asn Phe Ile Lys Val Phe Val
 115 120 125

 Pro Arg Asn Asp Glu Met Val Phe Val Cys Gly Thr Asn Ala Phe Asn
 130 135 140

 Pro Met Cys Arg Tyr Tyr Arg Val Ser Thr Leu Glu Tyr Asp Gly Glu
 145 150 155 160

 Glu Ile Ser Gly Leu Ala Arg Cys Pro Phe Asp Ala Arg Gln Thr Asn
 165 170 175

 Val Ala Leu Phe Ala Asp Gly Lys Leu Tyr Ser Ala Thr Val Ala Asp
 180 185 190

 Phe Leu Ala Ser Asp Ala Val Ile Tyr Arg Ser Met Gly Asp Gly Ser
 195 200 205

 Ala Leu Arg Thr Ile Lys Tyr Asp Ser Lys Trp Ile Lys Glu Pro His

210	215	220
Phe Leu His Ala Ile Glu Tyr Gly Asn Tyr Val Tyr Phe Phe Arg		
225	230	235
Glu Ile Ala Val Glu His Asn Asn Leu Gly Lys Ala Val Tyr Ser Arg		
245	250	255
Val Ala Arg Ile Cys Lys Asn Asp Met Gly Gly Ser Gln Arg Val Leu		
260	265	270
Glu Lys His Trp Thr Ser Phe Leu Lys Ala Arg Leu Asn Cys Ser Val		
275	280	285
Pro Gly Asp Ser Phe Phe Tyr Phe Asp Val Leu Gln Ser Ile Thr Asp		
290	295	300
Ile Ile Gln Ile Asn Gly Ile Pro Thr Val Val Gly Val Phe Thr Thr		
305	310	315
Gln Leu Asn Ser Ile Pro Gly Ser Ala Val Cys Ala Phe Ser Met Asp		
325	330	335
Asp Ile Glu Lys Val Phe Lys Gly Arg Phe Lys Glu Gln Lys Thr Pro		
340	345	350
Asp Ser Val Trp Thr Ala Val Pro Glu Asp Lys Val Pro Lys Pro Arg		
355	360	365
Pro Gly Cys Cys Ala Lys His Gly Leu Ala Glu Ala Tyr Lys Thr Ser		
370	375	380
Ile Asp Phe Pro Asp Glu Thr Leu Ser Phe Ile Lys Ser His Pro Leu		
385	390	395
Met Asp Ser Ala Val Pro Pro Ile Ala Asp Glu Pro Trp Phe Thr Lys		
405	410	415
Thr Arg Val Arg Tyr Arg Leu Thr Ala Ile Ser Val Asp His Ser Ala		
420	425	430
Gly Pro Tyr Gln Asn Tyr Thr Val Ile Phe Val Gly Ser Glu Ala Gly		
435	440	445
Met Val Leu Lys Val Leu Ala Lys Thr Ser Pro Phe Ser Leu Asn Asp		
450	455	460
Ser Val Leu Leu Glu Glu Ile Glu Ala Tyr Asn His Ala Lys Cys Ser		
465	470	475
Ala Glu Asn Glu Glu Asp Lys Val Ile Ser Leu Gln Leu Asp Lys		
485	490	495
Asp His His Ala Leu Tyr Val Ala Phe Ser Ser Cys Ile Ile Arg Ile		
500	505	510
Pro Leu Ser Arg Cys Glu Arg Tyr Gly Ser Cys Lys Lys Ser Cys Ile		

515	520	525
Ala Ser Arg Asp Pro Tyr Cys Gly Trp Leu Ser Gln Gly Ser Cys Gly		
530	535	540
Arg Val Thr Pro Gly Met Leu Leu Leu Thr Glu Asp Phe Phe Ala Phe		
545	550	560
His Asn His Ser Ala Glu Gly Tyr Glu Gln Asp Thr Glu Phe Gly Asn		
565	570	575
Thr Ala His Leu Gly Asp Cys His Glu Ile Leu Pro Thr Ser Thr Thr		
580	585	590
Pro Asp Tyr Lys Ile Phe Gly Gly Pro Thr Ser Gly Val Arg Trp Glu		
595	600	605
Val Gln Ser Gly Glu Ser Asn Gln Met Val His Met Asn Val Leu Ile		
610	615	620
Thr Cys Val Phe Ala Ala Phe Val Leu Gly Ala Phe Ile Ala Gly Val		
625	630	640
Ala Val Tyr Cys Tyr Arg Asp Met Phe Val Arg Lys Asn Arg Lys Ile		
645	650	655
His Lys Asp Ala Glu Ser Ala Gln Ser Cys Thr Asp Ser Ser Gly Ser		
660	665	670
Phe Ala Lys Leu Asn Gly Leu Phe Asp Ser Pro Val Lys Glu Tyr Gln		
675	680	685
Gln Asn Ile Asp Ser Pro Lys Leu Tyr Ser Asn Leu Leu Thr Ser Arg		
690	695	700
Lys Glu Leu Pro Pro Asn Gly Asp Ser Lys Ser Met Val Met Asp His		
705	710	720
Arg Gly Gln Pro Pro Glu Leu Ala Ala Leu Pro Thr Pro Glu Ser Thr		
725	730	735
Pro Val Leu His Gln Lys Thr Leu Gln Ala Met Lys Ser His Ser Glu		
740	745	750
Lys Ala His Gly His Gly Ala Ser Arg Lys Glu Thr Pro Gln Phe Phe		
755	760	765
Pro Ser Ser Pro Pro His Ser Pro Leu Ser His Gly His Ile Pro		
770	775	780
Ser Ala Ile Val Leu Pro Asn Ala Thr His Asp Tyr Asn Thr Ser Phe		
785	790	800
Ser Asn Ser Asn Ala His Lys Ala Glu Lys Lys Leu Gln Asn Ile Asp		
805	810	815
His Pro Leu Thr Lys Ser Ser Ser Lys Arg Asp His Arg Arg Ser Val		

820	825	830
Asp Ser Arg Asn Thr Leu Asn Asp Leu Leu Lys His Leu Asn Asp Pro		
835	840	845
Asn Ser Asn Pro Lys Ala Ile Met Gly Asp Ile Gln Met Ala His Gln		
850	855	860
Asn Leu Met Leu Asp Pro Met Gly Ser Met Ser Glu Val Pro Pro Lys		
865	870	875
Val Pro Asn Arg Glu Ala Ser Leu Tyr Ser Pro Pro Ser Thr Leu Pro		
885	890	895
Arg Asn Ser Pro Thr Lys Arg Val Asp Val Pro Thr Thr Pro Gly Val		
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Pro Met Thr Ser Leu Glu Arg Gln Arg Gly Tyr His Lys Asn Ser Ser		
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Gln Arg His Ser Ile Ser Ala Met Pro Lys Asn Leu Asn Ser Pro Asn		
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Gly Val Leu Leu Ser Arg Gln Pro Ser Met Asn Arg Gly Gly Tyr Met		
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Pro Thr Pro Thr Gly Ala Lys Val Asp Tyr Ile Gln Gly Thr Pro Val		
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Ser Val His Leu Gln Pro Ser Leu Ser Arg Gln Ser Ser Tyr Thr Ser		
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Asn Gly Thr Leu Pro Arg Thr Gly Leu Lys Arg Thr Pro Ser Leu Lys		
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Arg Pro Ser Gly Asn Glu Ser Gln His Arg Leu Asp Phe Gln Leu Met
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Leu Lys Ile Arg Asp Thr Leu Tyr Ile Ala Gly Arg Asp Gln Val Tyr
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Thr Val Asn Leu Asn Glu Met Pro Lys Thr Glu Val Ile Trp Gln Gln
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Lys Leu Thr Trp Arg Ser Arg Gln Gln Asp Arg Glu Asn Cys Ala Met
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<212> PRT

<213> Homo sapiens

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Arg	Pro	Ser	Gly	Asn	Glu	Ser	Gln	His	Arg	Leu	Asp	Phe	Gln	Leu	Met
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Leu	Lys	Ile	Arg	Asp	Thr	Leu	Tyr	Ile	Ala	Gly	Arg	Asp	Gln	Val	Tyr
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Lys	Leu	Thr	Trp	Arg	Ser	Arg	Gln	Gln	Asp	Arg	Glu	Asn	Cys	Ala	Met
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Lys	Gly	Lys	His	Lys	Asp	Glu	Cys	His	Asn	Phe	Ile	Lys	Val	Phe	Val
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 Pro Gly Cys Cys Ala Lys His Gly Leu Ala Glu Ala Tyr Lys Thr Ser
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Met Val Leu Lys Val Leu Ala Lys Thr Ser Pro Phe Ser Leu Asn Asp
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 Val Gln Asp Asp Pro Asn Thr Ser Asp Phe Thr Asp Pro Leu Ser Gly
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 Ile Pro Lys Gly Val Arg Trp Glu Val Gln Ser Gly Glu Ser Asn Gln
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 Met Val His Met Asn Val Leu Ile Thr Cys Val Phe Ala Ala Phe Val
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Tyr Ser Asn Leu Leu Thr Ser Arg Lys Glu Leu Pro Pro Asn Gly Asp
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Gln Ala Met Lys Ser His Ser Glu Lys Ala His Gly His Gly Ala Ser
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Arg Lys Glu Thr Pro Gln Phe Phe Pro Ser Ser Pro Pro His Ser
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Pro Leu Ser His Gly His Ile Pro Ser Ala Ile Val Leu Pro Asn Ala
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Thr His Asp Tyr Asn Thr Ser Phe Ser Asn Ser Asn Ala His Lys Ala
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Glu Lys Lys Leu Gln Asn Ile Asp His Pro Leu Thr Lys Ser Ser Ser
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Lys Arg Asp His Arg Arg Ser Val Asp Ser Arg Asn Thr Leu Asn Asp
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Arg Gly Tyr His Lys Asn Ser Ser Gln Arg His Ser Ile Ser Ala Met
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Pro Lys Asn Leu Asn Ser Pro Asn Gly Val Leu Leu Ser Arg Gln Pro
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Ser Met Asn Arg Gly Gly Tyr Met Pro Thr Pro Thr Gly Ala Lys Val
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Asp Tyr Ile Gln Gly Thr Pro Val Ser Val His Leu Gln Pro Ser Leu
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Ser Arg Gln Ser Ser Tyr Thr Ser Asn Gly Thr Leu Pro Arg Thr Gly
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<211> 698
<212> PRT
<213> Homo sapiens

<400> 22
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Trp Ser Ser Arg Ser Leu Gly Ala Arg Cys Arg Asn Ser Ile Ala Ser
35 40 45

Cys Pro Glu Glu Gln Pro His Val Gly Asn Tyr Arg Leu Leu Arg Thr
50 55 60

Ile Gly Lys Gly Asn Phe Ala Lys Val Lys Leu Ala Arg His Ile Leu
65 70 75 80

Thr Gly Arg Glu Val Ala Ile Lys Ile Ile Asp Lys Thr Gln Leu Asn
85 90 95

Pro Ser Ser Leu Gln Lys Leu Phe Arg Glu Val Arg Ile Met Lys Gly
100 105 110

Leu Asn His Pro Asn Ile Val Lys Leu Phe Glu Val Ile Glu Thr Glu
115 120 125

Lys Thr Leu Tyr Leu Val Met Glu Tyr Ala Ser Ala Gly Glu Pro Pro
130 135 140

Thr Leu Ser Ala Leu Pro Leu Cys His Leu Pro Leu Pro Leu His Leu
145 150 155 160

Thr Leu Thr Pro Leu Gly Leu Cys Pro Ala Gly Glu Val Phe Asp Tyr
165 170 175

Leu Val Ser His Gly Arg Met Lys Glu Lys Glu Ala Arg Ala Lys Phe
180 185 190

Arg Gln Ile Val Ser Ala Val His Tyr Cys His Gln Lys Asn Ile Val
195 200 205

His Arg Asp Leu Lys Ala Glu Asn Leu Leu Asp Ala Glu Ala Asn
210 215 220

Ile Lys Ile Ala Asp Phe Gly Phe Ser Asn Glu Phe Thr Leu Gly Ser
225 230 235 240

Lys Leu Asp Thr Phe Cys Gly Ser Pro Pro Tyr Ala Ala Pro Glu Leu
245 250 255

Phe Gln Gly Lys Lys Tyr Asp Gly Pro Glu Val Asp Ile Trp Ser Leu
260 265 270

Gly Val Ile Leu Tyr Thr Leu Val Ser Gly Ser Leu Pro Phe Asp Gly
 275 280 285
 His Asn Leu Lys Glu Leu Arg Glu Arg Val Leu Arg Gly Lys Tyr Arg
 290 295 300
 Val Pro Phe Tyr Met Ser Thr Asp Cys Glu Ser Ile Leu Arg Arg Phe
 305 310 315 320
 Leu Val Leu Asn Pro Ala Lys Arg Cys Thr Leu Glu Gln Ile Met Lys
 325 330 335
 Asp Lys Trp Ile Asn Ile Gly Tyr Glu Gly Glu Leu Lys Pro Tyr
 340 345 350
 Thr Glu Pro Glu Glu Asp Phe Gly Asp Thr Lys Arg Ile Glu Val Met
 355 360 365
 Val Gly Met Gly Tyr Thr Arg Glu Glu Ile Lys Glu Ser Leu Thr Ser
 370 375 380
 Gln Lys Tyr Asn Glu Val Thr Ala Thr Tyr Leu Leu Leu Gly Arg Lys
 385 390 395 400
 Leu Ser Pro Thr Ser Thr Gly Glu Ala Glu Leu Lys Glu Glu Arg Leu
 405 410 415
 Pro Gly Arg Lys Ala Ser Cys Ser Thr Ala Gly Ser Gly Ser Arg Gly
 420 425 430
 Leu Pro Pro Ser Ser Pro Met Val Ser Ser Ala His Asn Pro Asn Lys
 435 440 445
 Ala Glu Ile Pro Glu Arg Arg Lys Asp Ser Thr Pro Val Ser Asp Gln
 450 455 460
 Gly Trp Gly Met Met Thr Arg Arg Asn Thr Tyr Val Cys Thr Glu Arg
 465 470 475 480
 Pro Gly Ala Glu Arg Pro Ser Leu Leu Pro Asn Gly Lys Glu Asn Arg
 485 490 495
 Val Pro Pro Ala Ser Pro Ser Ser His Ser Leu Ala Pro Pro Ser Gly
 500 505 510
 Glu Arg Ser Arg Leu Ala Arg Gly Ser Thr Ile Arg Ser Thr Phe His
 515 520 525
 Gly Gly Gln Val Arg Asp Arg Arg Ala Gly Gly Gly Gly Gly Gly
 530 535 540
 Val Gln Asn Gly Pro Pro Ala Ser Pro Thr Leu Ala His Glu Ala Ala
 545 550 555 560
 Pro Leu Pro Ala Gly Arg Pro Arg Pro Thr Thr Asn Leu Phe Thr Lys
 565 570 575

Leu Thr Ser Lys Leu Thr Arg Ser Arg Leu Ser Cys His Leu Pro Trp
 580 585 590
 Asp Gln Thr Glu Thr Ala Pro Arg Leu Leu Arg Phe Pro Trp Ser Val
 595 600 605
 Lys Leu Thr Ser Ser Arg Pro Pro Glu Ala Leu Met Ala Ala Leu Arg
 610 615 620
 Gln Ala Thr Ala Ala Ala Arg Cys Arg Cys Arg Gln Pro Gln Pro Phe
 625 630 635 640
 Leu Leu Ala Cys Leu His Gly Gly Ala Gly Gly Pro Glu Pro Leu Ser
 645 650 655
 His Phe Glu Val Glu Val Cys Gln Leu Pro Arg Pro Gly Leu Arg Gly
 660 665 670
 Val Leu Phe Arg Arg Val Ala Gly Thr Ala Leu Ala Phe Arg Thr Leu
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 Val Thr Arg Ile Ser Asn Asp Leu Glu Leu
 690 695

<210> 23
 <211> 1549
 <212> DNA
 <213> Homo sapiens

<400> 23
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 gtccagccgc tcactgggtg cccgttgccg gaactccatc gcctcctgtc ccgaggagca 180
 gccccacgtg ggcaactacc gcctgctgag gaccattggg aagggaact ttgccaaagt 240
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 aaaccacccc aacatcgtga agctcttga ggtgatttag actgagaaga cgctgtacct 420
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 gaaggagaag gaagctcgag ccaagttccg acagattgtt tcggctgtc actattgtca 540
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 caacatcaag attgctgact ttggcttcag caacgagttc acgctggat cgaagctgga 660
 cacgttctgc gggagccccc catatgccgc cccggagctg tttcaggcga agaagtacga 720
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1549

<210> 24
<211> 508
<212> PRT
<213> Homo sapiens

<400> 24
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Thr His Gly Thr Leu Gly Ser Gly Arg Ser Ser Asp Lys Gly Pro Ser
20 25 30

Trp Ser Ser Arg Ser Leu Gly Ala Arg Cys Arg Asn Ser Ile Ala Ser
35 40 45

Cys Pro Glu Glu Gln Pro His Val Gly Asn Tyr Arg Leu Leu Arg Thr
50 55 60

Ile Gly Lys Gly Asn Phe Ala Lys Val Lys Leu Ala Arg His Ile Leu
65 70 75 80

Thr Gly Arg Glu Val Ala Ile Lys Ile Ile Asp Lys Thr Gln Leu Asn
85 90 95

Pro Ser Ser Leu Gln Lys Leu Phe Arg Glu Val Arg Ile Met Lys Gly
100 105 110

Leu Asn His Pro Asn Ile Val Lys Leu Phe Glu Val Ile Glu Thr Glu
115 120 125

Lys Thr Leu Tyr Leu Val Met Glu Tyr Ala Ser Ala Gly Glu Val Phe
130 135 140

Asp Tyr Leu Val Ser His Gly Arg Met Lys Glu Lys Glu Ala Arg Ala
145 150 155 160

Lys Phe Arg Gln Ile Val Ser Ala Val His Tyr Cys His Gln Lys Asn
165 170 175

Ile Val His Arg Asp Leu Lys Ala Glu Asn Leu Leu Asp Ala Glu
180 185 190

Ala Asn Ile Lys Ile Ala Asp Phe Gly Phe Ser Asn Glu Phe Thr Leu
195 200 205

Gly Ser Lys Leu Asp Thr Phe Cys Gly Ser Pro Pro Tyr Ala Ala Pro
210 215 220

Glu Leu Phe Gln Gly Lys Lys Tyr Asp Gly Pro Glu Val Asp Ile Trp
225 230 235 240

Ser Leu Gly Val Ile Leu Tyr Thr Leu Val Ser Gly Ser Leu Pro Phe
245 250 255

Asp Gly His Asn Leu Lys Glu Leu Arg Glu Arg Val Leu Arg Gly Lys
 260 265 270
 Tyr Arg Val Pro Phe Tyr Met Ser Thr Asp Cys Glu Ser Ile Leu Arg
 275 280 285
 Arg Phe Leu Val Leu Asn Pro Ala Lys Arg Cys Thr Leu Glu Gln Ile
 290 295 300
 Met Lys Asp Lys Trp Ile Asn Ile Gly Tyr Glu Gly Glu Glu Leu Lys
 305 310 315 320
 Pro Tyr Thr Glu Pro Glu Glu Asp Phe Gly Asp Thr Lys Arg Ile Glu
 325 330 335
 Val Met Val Gly Met Gly Tyr Thr Arg Glu Glu Ile Lys Glu Ser Leu
 340 345 350
 Thr Ser Gln Lys Tyr Asn Glu Val Thr Ala Gly Arg Pro Arg Pro Thr
 355 360 365
 Thr Asn Leu Phe Thr Lys Leu Thr Ser Lys Leu Thr Arg Arg Val Ala
 370 375 380
 Asp Glu Pro Glu Arg Ile Gly Gly Pro Glu Val Thr Ser Cys His Leu
 385 390 395 400
 Pro Trp Asp Gln Thr Glu Thr Ala Pro Arg Leu Leu Arg Phe Pro Trp
 405 410 415
 Ser Val Lys Leu Thr Ser Ser Arg Pro Pro Glu Ala Leu Met Ala Ala
 420 425 430
 Leu Arg Gln Ala Thr Ala Ala Arg Cys Arg Cys Arg Gln Pro Gln
 435 440 445
 Pro Phe Leu Leu Ala Cys Leu His Gly Gly Ala Gly Gly Pro Glu Pro
 450 455 460
 Leu Ser His Phe Glu Val Glu Val Cys Gln Leu Pro Arg Pro Gly Leu
 465 470 475 480
 Arg Gly Val Leu Phe Arg Arg Val Ala Gly Thr Ala Leu Ala Phe Arg
 485 490 495
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<210> 25
 <211> 4818
 <212> DNA
 <213> Homo sapiens

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<210> 26
<211> 1469
<212> PRT
<213> *Homo sapiens*

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<400> 26
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Pro Gly Phe Arg Ala Phe Leu Cys Pro Leu Ile Cys His Asn Gly Gly
35 40 45

Val Cys Val Lys Pro Asp Arg Cys Leu Cys Pro Pro Asp Phe Ala Gly
50 55 60

Lys Phe Cys Gln Leu His Ser Ser Gly Ala Arg Pro Pro Pro Ala Pro Ala
 65 70 75 80

Ile Pro Gly Leu Thr Arg Ser Val Tyr Thr Met Pro Leu Ala Asn His
 85 90 95

Arg Asp Asp Glu His Gly Val Ala Ser Met Val Ser Val His Val Glu
100 105 110

His Pro Gln Glu Ala Ser Val Val Val His Gln Val Glu Arg Val Ser
115 120 125

Gly Pro Trp Glu Glu Ala Asp Ala Glu Ala Val Ala Arg Ala Glu Ala
135 140

Ala Ala Arg Ala Glu Ala Ala Ala Pro Tyr Thr Val Leu Ala Gln Ser
 145 150 155 160
 Ala Pro Arg Glu Asp Gly Tyr Ser Asp Ala Ser Gly Phe Gly Tyr Cys
 165 170 175
 Phe Arg Glu Leu Arg Gly Gly Glu Cys Ala Ser Pro Leu Pro Gly Leu
 180 185 190
 Arg Thr Gln Glu Val Cys Cys Arg Gly Ala Gly Leu Ala Trp Gly Val
 195 200 205
 His Asp Cys Gln Leu Cys Ser Glu Arg Leu Gly Asn Ser Glu Arg Val
 210 215 220
 Ser Ala Pro Asp Gly Pro Cys Pro Thr Gly Phe Glu Arg Val Asn Gly
 225 230 235 240
 Ser Cys Glu Asp Val Asp Glu Cys Ala Thr Gly Gly Arg Cys Gln His
 245 250 255
 Gly Glu Cys Ala Asn Thr Arg Gly Gly Tyr Thr Cys Val Cys Pro Asp
 260 265 270
 Gly Phe Leu Leu Asp Ser Ser Arg Ser Ser Cys Ile Ser Gln His Val
 275 280 285
 Ile Ser Glu Ala Lys Gly Pro Cys Phe Arg Val Leu Arg Asp Gly Gly
 290 295 300
 Cys Ser Leu Pro Ile Leu Arg Asn Ile Thr Lys Gln Ile Cys Cys Cys
 305 310 315 320
 Ser Arg Val Gly Lys Ala Trp Gly Arg Gly Cys Gln Leu Cys Pro Pro
 325 330 335
 Phe Gly Ser Glu Gly Phe Arg Glu Ile Cys Pro Ala Gly Pro Gly Tyr
 340 345 350
 His Tyr Ser Ala Ser Asp Leu Arg Tyr Asn Thr Arg Pro Leu Gly Gln
 355 360 365
 Glu Pro Pro Arg Val Ser Leu Ser Gln Pro Arg Thr Leu Pro Ala Thr
 370 375 380
 Ser Arg Pro Ser Ala Gly Phe Leu Pro Thr His Arg Leu Glu Pro Arg
 385 390 395 400
 Pro Glu Pro Arg Pro Asp Pro Arg Pro Gly Pro Glu Phe Pro Leu Pro
 405 410 415
 Ser Ile Pro Ala Trp Thr Gly Pro Glu Ile Pro Glu Ser Gly Pro Ser
 420 425 430
 Ser Gly Met Cys Gln Arg Asn Pro Gln Val Cys Gly Pro Gly Arg Cys
 435 440 445

Ile Ser Arg Pro Ser Gly Tyr Thr Cys Ala Cys Asp Ser Gly Phe Arg
 450 455 460
 Leu Ser Pro Gln Gly Thr Arg Cys Ile Asp Val Asp Glu Cys Arg Arg
 465 470 475 480
 Val Pro Pro Pro Cys Ala Pro Gly Arg Cys Glu Asn Ser Pro Gly Ser
 485 490 495
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 Glu Cys Leu Asp Val Asp Glu Cys His Arg Val Pro Pro Pro Cys Asp
 515 520 525
 Leu Gly Arg Cys Glu Asn Thr Pro Gly Ser Phe Leu Cys Val Cys Pro
 530 535 540
 Ala Gly Tyr Gln Ala Ala Pro His Gly Ala Ser Cys Gln Asp Val Asp
 545 550 555 560
 Glu Cys Thr Gln Ser Pro Gly Leu Cys Gly Arg Gly Ala Cys Lys Asn
 565 570 575
 Leu Pro Gly Ser Phe Arg Cys Val Cys Pro Ala Gly Phe Arg Gly Ser
 580 585 590
 Ala Cys Glu Glu Asp Val Asp Glu Cys Ala Gln Glu Pro Pro Pro Cys
 595 600 605
 Gly Pro Gly Arg Cys Asp Asn Thr Ala Gly Ser Phe His Cys Ala Cys
 610 615 620
 Pro Ala Gly Phe Arg Ser Arg Gly Pro Gly Ala Pro Cys Gln Asp Val
 625 630 635 640
 Asp Glu Cys Ala Arg Ser Pro Pro Pro Cys Thr Tyr Gly Arg Cys Glu
 645 650 655
 Asn Thr Glu Gly Ser Phe Gln Cys Val Cys Pro Met Gly Phe Gln Pro
 660 665 670
 Asn Ala Ala Gly Ser Glu Cys Glu Asp Val Asp Glu Cys Glu Asn His
 675 680 685
 Leu Ala Cys Pro Gly Gln Glu Cys Val Asn Ser Pro Gly Ser Phe Gln
 690 695 700
 Cys Arg Ala Cys Pro Ser Gly His His Leu His Arg Gly Arg Cys Thr
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 Asp Val Asp Glu Cys Ser Ser Gly Ala Pro Pro Cys Gly Pro His Gly
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Arg Leu Asp Met Thr Arg Met Ala Cys Val Asp Ile Asn Glu Cys Asp
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Glu Ala Glu Ala Ala Ser Pro Leu Cys Val Asn Ala Arg Cys Leu Asn
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Lys Phe Cys Gln Leu His Ser Ser Gly Ala Arg Pro Pro Ala Pro Ala
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Ile Pro Gly Leu Thr Arg Ser Val Tyr Thr Met Pro Leu Ala Asn His
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His Pro Gln Glu Ala Ser Val Val Val His Gln Val Glu Arg Val Ser
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Gly Pro Trp Glu Glu Ala Asp Ala Glu Ala Val Ala Arg Ala Glu Ala
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Ala Pro Arg Glu Asp Gly Tyr Ser Asp Ala Ser Gly Phe Gly Tyr Cys
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Phe Arg Glu Leu Arg Gly Glu Cys Ala Ser Pro Leu Pro Gly Leu
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Arg Thr Gln Glu Val Cys Cys Arg Gly Ala Gly Leu Ala Trp Gly Val
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Ser Cys Glu Asp Val Asp Glu Cys Ala Thr Gly Gly Arg Cys Gln His
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Gly Glu Cys Ala Asn Thr Arg Gly Gly Tyr Thr Cys Val Cys Pro Asp
260 265 270

Gly Phe Leu Leu Asp Ser Ser Arg Ser Ser Cys Ile Ser Gln His Val
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Leu Ala Cys Pro Gly Gln Glu Cys Val Asn Ser Pro Gly Ser Phe Gln
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Cys Arg Ala Cys Pro Ser Gly His His Leu His Arg Gly Arg Cys Thr
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Asp Val Asp Glu Cys Ser Ser Gly Ala Pro Pro Cys Gly Pro His Gly
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His Cys Thr Asn Thr Glu Gly Ser Phe Arg Cys Ser Cys Ala Pro Gly
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 Lys Phe Cys Gln Leu His Ser Ser Gly Ala Arg Pro Pro Ala Pro Ala
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 His Pro Gln Glu Ala Ser Val Val His Gln Val Glu Arg Val Ser
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Lys	Phe	Cys	Gln	Leu	His	Ser	Ser	Gly	Ala	Arg	Pro	Pro	Ala	Pro	Ala
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 Val Asp Glu Cys Arg Glu Arg Gly Pro Ala Leu Cys Gly Ser Gln Arg
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 Cys Glu Asn Ser Pro Gly Ser Tyr Arg Cys Val Arg Asp Cys Asp Pro
 865 870 875 880
 Gly Tyr His Ala Gly Pro Glu Gly Thr Cys Asp Asp Val Asp Glu Cys
 885 890 895
 Arg Asn Arg Ser Phe Cys Gly Ala His Ala Val Cys Gln Asn Leu Pro
 900 905 910
 Gly Ser Phe Gln Cys Leu Cys Asp Gln Gly Tyr Glu Gly Ala Arg Asp
 915 920 925
 Gly Arg His Cys Val Asp Val Asn Glu Cys Glu Thr Leu Gln Gly Val
 930 935 940
 Cys Gly Ala Ala Leu Cys Glu Asn Val Glu Gly Ser Phe Leu Cys Val
 945 950 955 960
 Cys Pro Asn Ser Pro Glu Glu Phe Asp Pro Met Thr Gly Arg Cys Val
 965 970 975
 Pro Pro Arg Thr Ser Ala Asp Val Asp Glu Cys Gln Leu Phe Arg Asp
 980 985 990
 Gln Val Cys Lys Ser Gly Val Cys Val Asn Thr Ala Pro Gly Tyr Ser
 995 1000 1005
 Cys Tyr Cys Ser Asn Gly Tyr Tyr His Thr Gln Arg Leu Glu Cys
 1010 1015 1020
 Ile Asp Asn Asp Glu Cys Ala Asp Glu Glu Pro Ala Cys Glu Gly Gly
 1025 1030 1035 1040
 Arg Cys Val Asn Thr Val Gly Ser Tyr His Cys Thr Cys Glu Pro Pro
 1045 1050 1055

Leu Val Leu Asp Gly Ser Gln Arg Arg Cys Val Ser Asn Glu Ser Gln
 1060 1065 1070
 Ser Leu Asp Asp Asn Leu Gly Val Cys Trp Gln Glu Val Gly Ala Asp
 1075 1080 1085
 Leu Val Cys Ser His Pro Arg Leu Asp Arg Gln Ala Thr Tyr Thr Glu
 1090 1095 1100
 Cys Cys Cys Leu Tyr Gly Glu Ala Trp Gly Met Asp Cys Ala Leu Cys
 1105 1110 1115 1120
 Pro Ala Gln Asp Ser Asp Asp Phe Glu Ala Leu Cys Asn Val Leu Arg
 1125 1130 1135
 Pro Pro Ala Tyr Ser Pro Pro Arg Pro Gly Gly Phe Gly Leu Pro Tyr
 1140 1145 1150
 Glu Tyr Gly Pro Asp Leu Gly Pro Pro Tyr Gln Gly Leu Pro Tyr Gly
 1155 1160 1165
 Pro Glu Leu Tyr Pro Pro Ala Leu Pro Tyr Asp Pro Tyr Pro Pro
 1170 1175 1180
 Pro Pro Gly Pro Phe Ala Arg Arg Glu Ala Pro Tyr Gly Ala Pro Arg
 1185 1190 1195 1200
 Phe Asp Met Pro Asp Phe Glu Asp Asp Gly Gly Pro Tyr Gly Glu Ser
 1205 1210 1215
 Glu Ala Pro Ala Pro Pro Gly Pro Gly Thr Arg Trp Pro Tyr Arg Ser
 1220 1225 1230
 Arg Asp Thr Arg Arg Ser Phe Pro Glu Pro Glu Pro Pro Glu Gly
 1235 1240 1245
 Gly Ser Tyr Ala Gly Ser Leu Ala Glu Pro Tyr Glu Glu Leu Ala
 1250 1255 1260
 Glu Glu Cys Gly Ile Leu Asp Gly Cys Thr Asn Gly Arg Cys Val Arg
 1265 1270 1275 1280
 Val Pro Glu Gly Phe Thr Cys Arg Cys Phe Asp Gly Tyr Arg Leu Asp
 1285 1290 1295
 Met Thr Arg Met Ala Cys Val Asp Ile Asn Glu Cys Asp Glu Ala Glu
 1300 1305 1310
 Ala Ala Ser Pro Leu Cys Val Asn Ala Arg Cys Leu Asn Thr Asp Gly
 1315 1320 1325
 Ser Phe Arg Cys Ile Cys Arg Pro Gly Phe Ala Pro Thr His Gln Pro
 1330 1335 1340
 His His Cys Ala Pro Ala Arg Pro Arg Ala
 1345 1350

<210> 33
<211> 973
<212> DNA
<213> Homo sapiens

<400> 33
ggatgaacca gacttgaat agcagtggga ccgtggagtc agccctaaac tattccagag 60
ggagcacagt gcacacggcc tacctggtgc tgagctccct ggccatgttc acctgcctgt 120
gccccatggc aggcaacagc atggtgatct ggctgctagg ctttcgaatg cacaggaacc 180
ccttctgcat ctatattcctc aacctggcg cagccgaccc ccttccttc ttcagcatgg 240
cttccacgct cagcctggaa acccagcccc tggtaatac cactgacaag gtccacgagc 300
tggatgaagag actgatgtac ttgccttaca cagtgccctt gggctgctg acggccatca 360
gcacccagcg ctgtctctct gtcccttcc ctatctgggtt caagtgtcac cggccaggc 420
acctgtcagc ctgggtgtgt ggccctgtgt ggacgtctg ttcctgtatg aacgggttga 480
ccttccttcctt ctgcagcaag ttcttggaaat tcaatgaaga tcggtgcttc aggggtggaca 540
tgttccaggc cgccctcatc atgggggtct taaccccagt gatgactctg tccagcctga 600
ccctctttgt ctgggtgcgg aggagctccc agcagtggcg gcggcagccc acacggctgt 660
tcgtgggtgtt ctccttcctt gtccctgtgt tcctcatctg ttccctgtatg ctgagcatct 720
actgggttgt gctctactgg ttgagccgcg cgcccgagat gcaggtccctg tgcttcagct 780
tgtcacgcct ctcctcgtcc gtaaggcagca gcggccaaccc cgtcatctac ttccctgggtgg 840
gcagccggag gagccacagg ctgcccacca ggtccctggg gactgtgctc caacaggcgc 900
ttcgcgagga gcccggagctg gaaggtgggg agacggccac cgtggcacc aatgagatgg 960
gggcttgaga gcc 973

<210> 34
<211> 321
<212> PRT
<213> Homo sapiens

<400> 34
Met Asn Gln Thr Leu Asn Ser Ser Gly Thr Val Glu Ser Ala Leu Asn
1 5 10 15
Tyr Ser Arg Gly Ser Thr Val His Thr Ala Tyr Leu Val Leu Ser Ser
20 25 30
Leu Ala Met Phe Thr Cys Leu Cys Gly Met Ala Gly Asn Ser Met Val
35 40 45
Ile Trp Leu Leu Gly Phe Arg Met His Arg Asn Pro Phe Cys Ile Tyr
50 55 60
Ile Leu Asn Leu Ala Ala Ala Asp Leu Leu Phe Leu Phe Ser Met Ala
65 70 75 80
Ser Thr Leu Ser Leu Glu Thr Gln Pro Leu Val Asn Thr Thr Asp Lys
85 90 95
Val His Glu Leu Met Lys Arg Leu Met Tyr Phe Ala Tyr Thr Val Gly
100 105 110
Leu Ser Leu Leu Thr Ala Ile Ser Thr Gln Arg Cys Leu Ser Val Leu
115 120 125
Phe Pro Ile Trp Phe Lys Cys His Arg Pro Arg His Leu Ser Ala Trp

130	135	140
Val Cys Gly Leu Leu Trp Thr Leu Cys Leu Leu Met Asn Gly Leu Thr		
145	150	155
Ser Ser Phe Cys Ser Lys Phe Leu Lys Phe Asn Glu Asp Arg Cys Phe		
165	170	175
Arg Val Asp Met Val Gln Ala Ala Leu Ile Met Gly Val Leu Thr Pro		
180	185	190
Val Met Thr Leu Ser Ser Leu Thr Leu Phe Val Trp Val Arg Arg Ser		
195	200	205
Ser Gln Gln Trp Arg Arg Gln Pro Thr Arg Leu Phe Val Val Val Leu		
210	215	220
Ala Ser Val Leu Val Phe Leu Ile Cys Ser Leu Pro Leu Ser Ile Tyr		
225	230	240
Trp Phe Val Leu Tyr Trp Leu Ser Pro Pro Pro Glu Met Gln Val Leu		
245	250	255
Cys Phe Ser Leu Ser Arg Leu Ser Ser Ser Val Ser Ser Ala Asn		
260	265	270
Pro Val Ile Tyr Phe Leu Val Gly Ser Arg Arg Ser His Arg Leu Pro		
275	280	285
Thr Arg Ser Leu Gly Thr Val Leu Gln Gln Ala Leu Arg Glu Glu Pro		
290	295	300
Glu Leu Glu Gly Gly Glu Thr Pro Thr Val Gly Thr Asn Glu Met Gly		
305	310	320

Ala

<210> 35
<211> 671
<212> DNA
<213> Homo sapiens

<400> 35
atgcgaagtc actcttaccc tctgatgataa taatgggtat aattttcttg gtgcctctgt 60
tctggggaaa tgaggttaat gatgaaggcag tgatgtcaac tttagaacac ttgcattgtgg 120
actaccctca gaatgacgtt cccgttccctg caaggtaactg caaccacatg atcatacaaa 180
gagtttatcag ggaacctgac cacacttcta aaaaggagca tgtcttcatc catgagaggc 240
ctcgaaaaat caatgttatt tgcatttctc ccaagaagggt tgcttgccaa aaccttcgg 300
ccatttctg ctccatcaggat gagacaaaagt tcaaaatgac agtctgtcaag ctcattgaag 360
gcacaagata ccctgcctgc aggttaccact attcccccac agaggggtt gttcttgtca 420
cttgcgtatca cttgaggcca gatagttcc ttggctatgt taaataactc aagatcagct 480
cccgagtcg agatcttcc tctcaatggc attggagctg gctgtgcctg aggccagacct 540
ggaccgtgga catggggcaa tgccttgaac ggaaggggaa gccactggta attaattttat 600
ccttcctgtt ttgcgggtt gggattgttt tattctgctt caataaaata atctttactg 660
aattaaaaaaaaa a 671

<210> 36
 <211> 154
 <212> PRT
 <213> Homo sapiens

<400> 36
 Ala Lys Ser Leu Leu Pro Leu Met Ile Ile Met Val Ile Ile Phe Leu
 1 5 10 15

Val Leu Leu Phe Trp Glu Asn Glu Val Asn Asp Glu Ala Val Met Ser
 20 25 30

Thr Leu Glu His Leu His Val Asp Tyr Pro Gln Asn Asp Val Pro Val
 35 40 45

Pro Ala Arg Tyr Cys Asn His Met Ile Ile Gln Arg Val Ile Arg Glu
 50 55 60

Pro Asp His Thr Cys Lys Lys Glu His Val Phe Ile His Glu Arg Pro
 65 70 75 80

Arg Lys Ile Asn Gly Ile Cys Ile Ser Pro Lys Lys Val Ala Cys Gln
 85 90 95

Asn Leu Ser Ala Ile Phe Cys Phe Gln Ser Glu Thr Lys Phe Lys Met
 100 105 110

Thr Val Cys Gln Leu Ile Glu Gly Thr Arg Tyr Pro Ala Cys Arg Tyr
 115 120 125

His Tyr Ser Pro Thr Glu Gly Phe Val Leu Val Thr Cys Asp Asp Leu
 130 135 140

Arg Pro Asp Ser Phe Leu Gly Tyr Val Lys
 145 150

<210> 37
 <211> 1476
 <212> DNA
 <213> Homo sapiens

<400> 37
 caggtgcaaa ccagccccag gctccatggc ttcaagaagg tcgaagttca agggaaagcac 60
 caaggctccc ttgtgggtct gaaaaatctgc attggtaat gctttaggct ttttacttc 120
 ttcatgcaaa gtttctttg catcgatcc catcaaaaata gtgagagccc agaggcagta 180
 catgtttgat gagaacggtg aacagtactt ggactgcattt aacaatgttgc cggtggaca 240
 ctgtcaccca ggagtggtca aagctgcctt gaaacagatg gaactgctaa atacaaattc 300
 tcgattcctc cacgacaaca ttgttgatgta tgccaaacgc ctttcagcaa ctctgcccga 360
 gaaaactctt gtttggattt ttacaaaattc agggtccgaa gccaacgact tagccttacg 420
 .cctggctcgg cagttcagag gccaccagga tgtgatcaactt cttgacgctt accatggtca 480
 ccttatcatcc ttaattgaga ttagccata taagtttcag aaaggaaaaag atgtcaaaaa 540
 agaatttcta catgtggcac caactccaga tacttacaga gggaaatata gagaagacca 600
 tgcagactca gccagtgcattt atgcagatga agtgaagaaaa atcattgaag atgctcataa 660
 cagtggaaagg aagttgctg ctttattgc tgaatccatg cagagttgtg gcggacaaat 720

aattcctcca gcaggctact tccagaaaagt ggcagagtat gtacacgggt caggggggtgt 780
 gtttatagct gatgaagttc aagtgggctt tggcagagtt gggaaacatt tctggagctt 840
 ccagatgtat ggtgaagact ttgttccaga catcgtcaca atgggaaaac cgatggcaa 900
 cggccaccccg gtggcatgtg tggttaacaac caaagaaatt gcagaagct tcagcagctc 960
 tgggatggaa tatttaata cgtatggagg aaatccagta tcttgtgctg ttgggttggc 1020
 tgcctggat ataattgaaa atgaagacct tcaaggaaat gccaagagag taggaatta 1080
 tctcactgag ttactgaaaa aacagaaggc taaacacact ttgataggag atattagggg 1140
 cattggcctt tttattggaa ttgatttagt gaaggaccat ctgaaaagga cccctgatat 1200
 gtatttagct ttggggacaa ttttggttct ggagaaagaa aaacgagtgc ttctcagtgc 1260
 cgatggacct catagaaaatg tacttaaaat aaaaccacct atgtgcttca ctgaagaaga 1320
 tgcaaagttc atggtgacc aacttgatag gattctaaca gttgggtcca tggatctta 1380
 agatgtcttc ttgttccctc tcccaaacc acccctcaaa ccctggtcta gtcataatga 1440
 gcatatgcat cttgttattc atgatgaaag tgaggc 1476

<210> 38
 <211> 451
 <212> PRT
 <213> Homo sapiens

<400> 38
 Met Ala Ser Arg Arg Ser Lys Phe Lys Gly Ser Thr Lys Ala Pro Leu
 1 5 10 15
 Trp Val Trp Lys Ser Ala Leu Val Asn Ala Leu Gly Phe Phe Thr Ser
 20 25 30
 Ser Cys Lys Val Phe Phe Ala Ser Asp Pro Ile Lys Ile Val Arg Ala
 35 40 45
 Gln Arg Gln Tyr Met Phe Asp Glu Asn Gly Glu Gln Tyr Leu Asp Cys
 50 55 60
 Ile Asn Asn Val Ala Val Gly His Cys His Pro Gly Val Val Lys Ala
 65 70 75 80
 Ala Leu Lys Gln Met Glu Leu Leu Asn Thr Asn Ser Arg Phe Leu His
 85 90 95
 Asp Asn Ile Val Glu Tyr Ala Lys Arg Leu Ser Ala Thr Leu Pro Glu
 100 105 110
 Lys Leu Ser Val Cys Tyr Phe Thr Asn Ser Gly Ser Glu Ala Asn Asp
 115 120 125
 Leu Ala Leu Arg Leu Ala Arg Gln Phe Arg Gly His Gln Asp Val Ile
 130 135 140
 Thr Leu Asp Ala Tyr His Gly His Leu Ser Ser Leu Ile Glu Ile Ser
 145 150 155 160
 Pro Tyr Lys Phe Gln Lys Gly Lys Asp Val Lys Lys Glu Phe Val His
 165 170 175
 Val Ala Pro Thr Pro Asp Thr Tyr Arg Gly Lys Tyr Arg Glu Asp His
 180 185 190

Ala Asp Ser Ala Ser Ala Tyr Ala Asp Glu Val Lys Lys Ile Ile Glu
 195 200 205
 Asp Ala His Asn Ser Gly Arg Lys Val Ala Ala Phe Ile Ala Glu Ser
 210 215 220
 Met Gln Ser Cys Gly Gly Gln Ile Ile Pro Pro Ala Gly Tyr Phe Gln
 225 230 235 240
 Lys Val Ala Glu Tyr Val His Gly Ala Gly Gly Val Phe Ile Ala Asp
 245 250 255
 Glu Val Gln Val Gly Phe Gly Arg Val Gly Lys His Phe Trp Ser Phe
 260 265 270
 Gln Met Tyr Gly Glu Asp Phe Val Pro Asp Ile Val Thr Met Gly Lys
 275 280 285
 Pro Met Gly Asn Gly His Pro Val Ala Cys Val Val Thr Thr Lys Glu
 290 295 300
 Ile Ala Glu Ala Phe Ser Ser Ser Gly Met Glu Tyr Phe Asn Thr Tyr
 305 310 315 320
 Gly Gly Asn Pro Val Ser Cys Ala Val Gly Leu Ala Val Leu Asp Ile
 325 330 335
 Ile Glu Asn Glu Asp Leu Gln Gly Asn Ala Lys Arg Val Gly Asn Tyr
 340 345 350
 Leu Thr Glu Leu Leu Lys Lys Gln Lys Ala Lys His Thr Leu Ile Gly
 355 360 365
 Asp Ile Arg Gly Ile Gly Leu Phe Ile Gly Ile Asp Leu Val Lys Asp
 370 375 380
 His Leu Lys Arg Thr Pro Asp Met Tyr Leu Ala Leu Gly Thr Ile Leu
 385 390 395 400
 Val Leu Glu Lys Glu Lys Arg Val Leu Leu Ser Ala Asp Gly Pro His
 405 410 415
 Arg Asn Val Leu Lys Ile Lys Pro Pro Met Cys Phe Thr Glu Glu Asp
 420 425 430
 Ala Lys Phe Met Val Asp Gln Leu Asp Arg Ile Leu Thr Gly Gly Ser
 435 440 445
 Met Asp Leu
 450

<210> 39
 <211> 3350
 <212> DNA
 <213> Homo sapiens

<400> 39

cgccccattgg ctcctcagcc aagcacgtac accaaatgtc tgaacctgcg gttccctctcg 60
tactgagcag gattaccatg gcaacaacac atcatcaga gggtaaaact aacctgtctc 120
acgacggctc aaaccaggc agcctcgcc gccggcaag tagctccgag cggctgctc 180
ccggttgcct cgaagaagac agggggcgcc gcgcctcgct tgctccgdc ctgagccatg 240
cccagcagcc ctgtgttaacc accgagtccc ggccggagcc gaccgaccca gtgtgcggc 300
tcttcggcc gagctgagct ttctgtcagc caactccctc tgccccagcc ggccccggc 360
caccatgccc cgggcgactg cacttgggc cctggtgtca ctgctgctgc tgctgccgt 420
gcctcgccg gcccgggac tcggggagcg cccggacgac accgcagact actcagagct 480
ggacggcgag gagggcacgg agcagcagct ggagcattac cacgaccctt gcaaagccgc 540
tgtctttgg ggagacattt ccttagatga agatgactt aagctgttc acattgacaa 600
agccagagac tggaccaaggc agacagtggg ggcaacacgga cacagcacag gtgggcttga 660
agagcaggca tctgagagca gcccagacac cacagccatg gacactggca ccaaggaagc 720
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agccaagacc ttctctcccc gggtccgaag agccacaacc tcaaggacag agaggatatg 840
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agaccaacat gtcaccatca tcagggaaaa catccagcca ggtcaggagt ataatttctt 1200
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gtcccgatggc tctatcaata aagccggcct tgcaaggccat ttttcaagg aggtggatga 2100
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caagtgtgcc tgtgaccctg gctacgagct gcccggcgt aagaagatgt gtgaagtggc 2220
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gtatcccaca aacaaaaact gtgtctggca ggtggggcccccactcgat accggatctc 2340
ccttcagttt gaagtgtttt aactggaaagg caatgacgtc tgtaagtacg actttgtaga 2400
ggtgccgcgc ggcctgtccc cccgcacccaa gctgcacggc aggttctgc gctctgagac 2460
gccggaaagtc atcacctcgca agagcaacaa catgcgcgtg gagttcaagt ccgacaacac 2520
cgctctccaag cgccggctca gggcccaactt cttctcagat aaggacgagt gtgccaagga 2580
caacggcggg tgtcagcatg agtgcgtcaa caccttcggg agctacctgt gcaggtgcag 2640
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gatcagcagt gtggagggga ccctggcgag ccccaactgg cctgacaaat accccagccg 2760
gagggaggtgt acctgaaaca tctttcgac tgcaaggccac agagtgaaac tcacctttaa 2820
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gccggacacgc ctggccccca ttctggccg tttctgcggc agcaagaaac cagacccac 2940
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ctattccac gcccagtttgg gggacaacaa ctacccgagc gaggcccgt gtgactgggt 3060
gatcgtggca gaggacggct acggcggtga gctgacattc cggaccccttgc aggttgcgg 3120
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caggctcgcc cgcttctgtg gctctggggc attagaagaa atctactctg caggtgattc 3240
cctgtatgatt cgattccgca cagatgacac catcaacaag aaaggcttc atgcccgtata 3300
caccagcacc aagttccagg atggcctgca catgaagaaa tagtgcgtat 3350

<210> 40
 <211> 992
 <212> PRT
 <213> Homo sapiens

<400> 40
 Met Pro Arg Ala Thr Ala Leu Gly Ala Leu Val Ser Leu Leu Leu
 1 5 10 15

Leu Pro Leu Pro Arg Gly Ala Gly Gly Leu Gly Glu Arg Pro Asp Ala
 20 25 30

Thr Ala Asp Tyr Ser Glu Leu Asp Gly Glu Glu Gly Thr Glu Gln Gln
 35 40 45

Leu Glu His Tyr His Asp Pro Cys Lys Ala Ala Val Phe Trp Gly Asp
 50 55 60

Ile Ala Leu Asp Glu Asp Asp Leu Lys Leu Phe His Ile Asp Lys Ala
 65 70 75 80

Arg Asp Trp Thr Lys Gln Thr Val Gly Ala Thr Gly His Ser Thr Gly
 85 90 95

Gly Leu Glu Glu Gln Ala Ser Glu Ser Ser Pro Asp Thr Thr Ala Met
 100 105 110

Asp Thr Gly Thr Lys Glu Ala Gly Lys Asp Gly Arg Glu Asn Thr Thr
 115 120 125

Leu Leu His Ser Pro Gly Thr Leu His Ala Ala Ala Lys Thr Phe Ser
 130 135 140

Pro Arg Val Arg Arg Ala Thr Thr Ser Arg Thr Glu Arg Ile Trp Pro
 145 150 155 160

Gly Gly Val Ile Pro Tyr Val Ile Gly Gly Asn Phe Thr Gly Ser Gln
 165 170 175

Arg Ala Ile Phe Lys Gln Ala Met Arg His Trp Glu Lys His Thr Cys
 180 185 190

Val Thr Phe Ile Glu Arg Thr Asp Glu Glu Ser Phe Ile Val Phe Ser
 195 200 205

Tyr Arg Thr Cys Gly Cys Cys Ser Tyr Val Gly Arg Arg Gly Gly Gly
 210 215 220

Pro Gln Ala Ile Ser Ile Gly Lys Asn Cys Asp Lys Phe Gly Ile Val
 225 230 235 240

Ala His Glu Leu Gly His Val Val Gly Phe Trp His Glu His Thr Arg
 245 250 255

Pro Asp Arg Asp Gln His Val Thr Ile Ile Arg Glu Asn Ile Gln Pro

260	265	270
Gly Gln Glu Tyr Asn Phe Leu Lys Met Glu Ala Gly Glu Val Ser Ser		
275	280	285
Leu Gly Glu Thr Tyr Asp Phe Asp Ser Ile Met His Tyr Ala Arg Asn		
290	295	300
Thr Phe Ser Arg Gly Val Phe Leu Asp Thr Ile Leu Pro Arg Gln Asp		
305	310	315
Asp Asn Gly Val Arg Pro Thr Ile Gly Gln Arg Val Arg Leu Ser Gln		
325	330	335
Gly Asp Ile Ala Gln Ala Arg Lys Leu Tyr Lys Cys Pro Ala Cys Gly		
340	345	350
Glu Thr Leu Gln Asp Thr Thr Gly Asn Phe Ser Ala Pro Gly Phe Pro		
355	360	365
Asn Gly Tyr Pro Ser Tyr Ser His Cys Val Trp Arg Ile Ser Val Thr		
370	375	380
Pro Gly Glu Lys Ile Val Leu Asn Phe Thr Ser Met Asp Leu Phe Lys		
385	390	395
Ser Arg Leu Cys Trp Tyr Asp Tyr Val Glu Val Arg Asp Gly Tyr Trp		
405	410	415
Arg Lys Ala Pro Leu Leu Gly Arg Phe Cys Gly Asp Lys Ile Pro Glu		
420	425	430
Pro Leu Val Ser Thr Asp Ser Arg Leu Trp Val Glu Phe Arg Ser Ser		
435	440	445
Ser Asn Ile Leu Gly Lys Gly Phe Phe Ala Ala Tyr Glu Ala Thr Cys		
450	455	460
Gly Gly Asp Met Asn Lys Asp Ala Gly Gln Ile Gln Ser Pro Asn Tyr		
465	470	480
Pro Asp Asp Tyr Arg Pro Ser Lys Glu Cys Val Trp Arg Ile Thr Val		
485	490	495
Ser Glu Gly Phe His Val Gly Leu Thr Phe Gln Ala Phe Glu Ile Glu		
500	505	510
Arg His Asp Ser Cys Ala Tyr Asp Tyr Leu Glu Val Arg Asp Gly Pro		
515	520	525
Thr Glu Glu Ser Ala Leu Ile Gly His Phe Cys Gly Tyr Glu Lys Pro		
530	535	540
Glu Asp Val Lys Ser Ser Ser Asn Arg Leu Trp Met Lys Phe Val Ser		
545	550	560
Asp Gly Ser Ile Asn Lys Ala Gly Phe Ala Ala Asn Phe Phe Lys Glu		

565	570	575
Val Asp Glu Cys Ser Trp Pro Asp His Gly Gly Cys Glu His Arg Cys		
580	585	590
Val Asn Thr Leu Gly Ser Tyr Lys Cys Ala Cys Asp Pro Gly Tyr Glu		
595	600	605
Leu Ala Ala Asp Lys Lys Met Cys Glu Val Ala Cys Gly Gly Phe Ile		
610	615	620
Thr Lys Leu Asn Gly Thr Ile Thr Ser Pro Gly Trp Pro Lys Glu Tyr		
625	630	640
Pro Thr Asn Lys Asn Cys Val Trp Gln Val Val Ala Pro Thr Gln Tyr		
645	650	655
Arg Ile Ser Leu Gln Phe Glu Val Phe Glu Leu Glu Gly Asn Asp Val		
660	665	670
Cys Lys Tyr Asp Phe Val Glu Val Arg Ser Gly Leu Ser Pro Asp Ala		
675	680	685
Lys Leu His Gly Arg Phe Cys Gly Ser Glu Thr Pro Glu Val Ile Thr		
690	695	700
Ser Gln Ser Asn Asn Met Arg Val Glu Phe Lys Ser Asp Asn Thr Val		
705	710	720
Ser Lys Arg Gly Phe Arg Ala His Phe Ser Asp Lys Asp Glu Cys		
725	730	735
Ala Lys Asp Asn Gly Gly Cys Gln His Glu Cys Val Asn Thr Phe Gly		
740	745	750
Ser Tyr Leu Cys Arg Cys Arg Asn Gly Tyr Trp Leu His Glu Asn Gly		
755	760	765
His Asp Cys Lys Glu Ala Gly Cys Ala His Lys Ile Ser Ser Val Glu		
770	775	780
Gly Thr Leu Ala Ser Pro Asn Trp Pro Asp Lys Tyr Pro Ser Arg Arg		
785	790	800
Glu Cys Thr Trp Asn Ile Ser Ser Thr Ala Gly His Arg Val Lys Leu		
805	810	815
Thr Phe Asn Glu Phe Glu Ile Glu Gln His Gln Glu Cys Ala Tyr Asp		
820	825	830
His Leu Glu Met Tyr Asp Gly Pro Asp Ser Leu Ala Pro Ile Leu Gly		
835	840	845
Arg Phe Cys Gly Ser Lys Lys Pro Asp Pro Thr Val Ala Ser Gly Ser		
850	855	860
Lys Cys Gly Gly Arg Leu Lys Ala Glu Val Gln Thr Lys Glu Leu Tyr		

865	870	875	880
Ser His Ala Gln Phe Gly Asp Asn Asn Tyr Pro Ser Glu Ala Arg Cys			
885	890	895	
Asp Trp Val Ile Val Ala Glu Asp Gly Tyr Gly Val Glu Leu Thr Phe			
900	905	910	
Arg Thr Phe Glu Val Glu Glu Ala Asp Cys Gly Tyr Asp Tyr Met			
915	920	925	
Glu Ala Tyr Asp Gly Tyr Asp Ser Ser Ala Pro Arg Leu Gly Arg Phe			
930	935	940	
Cys Gly Ser Gly Pro Leu Glu Glu Ile Tyr Ser Ala Gly Asp Ser Leu			
945	950	955	960
Met Ile Arg Phe Arg Thr Asp Asp Thr Ile Asn Lys Lys Gly Phe His			
965	970	975	
Ala Arg Tyr Thr Ser Thr Lys Phe Gln Asp Gly Leu His Met Lys Lys			
980	985	990	

<210> 41
 <211> 3146
 <212> DNA
 <213> Homo sapiens

<400> 41
 gcagcctcg ccggccggca agtagctccg agcggctgct tcccggttgc ctcgacgaag 60
 acagggggccc cgcgcgtccg cttgctccgc gcctgagcca tgccccagcag ccctgtgtaa 120
 ccaccgagtc ccggccggag ccgaccgacc cagtgtgcgc cgtctttcgg ccgagctgag 180
 cttegtcga cgcaactccc tctgccccag ccggcccccgc gccaccatgc cccggggcgcac 240
 tgcacttggg gccccttgtt cactgctgct gctgctgccc ctgcctcgcg ggcggccgggg 300
 actcgaaaaa cggccggacg ccaccgcaga ctactcagag ctggacggcg aggagggcac 360
 ggagcagcag ctggagcatt accacgaccc ttgcaaagcc gctgtcttt gggagacat 420
 tgccttagat gaagatgact tgaagcttt tcacattgac aaagccagag actggaccaa 480
 gcagacagtg gggcaaacag gacacagcac aggtgggctt gaagagcagg catctgagag 540
 cagccccagac accacagcca tggacactgg caccaaggaa gctggaaaagg ggagccagag 600
 ggcattttt aagcaggcca tgagacactg ggagaagcac acctgtgtga ctttcataga 660
 aaggacggat gaggaaagct ttattgtatt cagttacaga acctgtggct tttgtccta 720
 ttttggggcgc cgaggaggag gcccacaggc catatccatt gggaaaact gtgacaagtt 780
 tggcatttgt gtcacgagc tggccatgt gttgggttt tggcatgaac acacccggcc 840
 agacagagac caacatgtca ccatcatcag gaaaaacatc cagccaggc aggagtataa 900
 tttcttaaaa atgaaagctg gggaaagttag ctctctggaa gagacatacg acttgacag 960
 catcatgcac tacgccccga acacccctc aagaggagtt ttttttagaca ccatcccttcc 1020
 ccgtcaagat gacaatggcg tcaggccaaac cattggccag cgcgtgcggc tcagtcaggg 1080
 agacatagct caagccccga agctgtacaa atgcccaggc cctacttgtt cttttgttag 1140
 ccagaaaaca tcaatctgct tgctacactt ctcaccaacc ttttccggagg gctttggctg 1200
 gcaaaggccg tttggggaga ccctgcaggc cacaacggga aactttctg cacctggttt 1260
 cccaaatggg taccatctt actcccactg cgtctggagg atctcggtca ccccaagggg 1320
 aaagatcgta taaaacttca catccatgga ttttttaaa agccgactgt gctgttatga 1380
 ttacgtggag gtccgggatg gttactggag aaaagcccccc cttttggca ggtttgtgg 1440

<210> 42
<211> 970
<212> PRT
<213> *Homo sapiens*

<400> 42
Met Pro Arg Ala Thr Ala Leu Gly Ala Leu Val Ser Leu Leu Leu Leu
1 5 10 15

Leu Pro Leu Pro Arg Gly Ala Gly Gly Leu Gly Glu Arg Pro Asp Ala
 20 25 30

Thr Ala Asp Tyr Ser Glu Leu Asp Gly Glu Glu Gly Thr Glu Gln Gln
 35 40 45

Leu Glu His Tyr His Asp Pro Cys Lys Ala Ala Val Phe Trp Gly Asp
50 55 60

Ile Ala Leu Asp Glu Asp Asp Leu Lys Leu Phe His Ile Asp Lys Ala
65 70 75 80

Arg Asp Trp Thr Lys Gln Thr Val Gly Ala Thr Gly His Ser Thr Gly
85 90 95

Gly Leu Glu Glu Gln Ala Ser Glu Ser Ser Pro Asp Thr Thr Ala Met
100 105 110

Asp Thr Gly Thr Lys Glu Ala Gly Lys Ser Gln Arg Ala Ile Phe
 115 120 125
 Lys Gln Ala Met Arg His Trp Glu Lys His Thr Cys Val Thr Phe Ile
 130 135 140
 Glu Arg Thr Asp Glu Glu Ser Phe Ile Val Phe Ser Tyr Arg Thr Cys
 145 150 155 160
 Gly Cys Cys Ser Tyr Val Gly Arg Arg Gly Gly Pro Gln Ala Ile
 165 170 175
 Ser Ile Gly Lys Asn Cys Asp Lys Phe Gly Ile Val Ala His Glu Leu
 180 185 190
 Gly His Val Val Gly Phe Trp His Glu His Thr Arg Pro Asp Arg Asp
 195 200 205
 Gln His Val Thr Ile Ile Arg Glu Asn Ile Gln Pro Gly Gln Glu Tyr
 210 215 220
 Asn Phe Leu Lys Met Glu Ala Gly Glu Val Ser Ser Leu Gly Glu Thr
 225 230 235 240
 Tyr Asp Phe Asp Ser Ile Met His Tyr Ala Arg Asn Thr Phe Ser Arg
 245 250 255
 Gly Val Phe Leu Asp Thr Ile Leu Pro Arg Gln Asp Asp Asn Gly Val
 260 265 270
 Arg Pro Thr Ile Gly Gln Arg Val Arg Leu Ser Gln Gly Asp Ile Ala
 275 280 285
 Gln Ala Arg Lys Leu Tyr Lys Cys Pro Gly Pro Thr Cys Ala Phe Val
 290 295 300
 Ser Gln Lys Thr Ser Ile Cys Leu Leu His Phe Ser Pro Thr Cys Ser
 305 310 315 320
 Glu Gly Phe Gly Trp Gln Arg Ala Cys Gly Glu Thr Leu Gln Asp Thr
 325 330 335
 Thr Gly Asn Phe Ser Ala Pro Gly Phe Pro Asn Gly Tyr Pro Ser Tyr
 340 345 350
 Ser His Cys Val Trp Arg Ile Ser Val Thr Pro Gly Glu Lys Ile Val
 355 360 365
 Leu Asn Phe Thr Ser Met Asp Leu Phe Lys Ser Arg Leu Cys Trp Tyr
 370 375 380
 Asp Tyr Val Glu Val Arg Asp Gly Tyr Trp Arg Lys Ala Pro Leu Leu
 385 390 395 400
 Gly Arg Phe Cys Gly Asp Lys Ile Pro Glu Pro Leu Val Ser Thr Asp
 405 410 415

Ser Arg Leu Trp Val Glu Phe Arg Ser Ser Ser Asn Ile Leu Gly Lys
 420 425 430
 Gly Phe Phe Ala Ala Tyr Glu Ala Thr Cys Gly Gly Asp Met Asn Lys
 435 440 445
 Asp Ala Gly Gln Ile Gln Ser Pro Asn Tyr Pro Asp Asp Tyr Arg Pro
 450 455 460
 Ser Lys Glu Cys Val Trp Arg Ile Thr Val Ser Glu Gly Phe His Val
 465 470 475 480
 Gly Leu Thr Phe Gln Ala Phe Glu Ile Glu Arg His Asp Ser Cys Ala
 485 490 495
 Tyr Asp Tyr Leu Glu Val Arg Asp Gly Pro Thr Glu Glu Ser Ala Leu
 500 505 510
 Ile Gly His Phe Cys Gly Tyr Glu Lys Pro Glu Asp Val Lys Ser Ser
 515 520 525
 Ser Asn Arg Leu Trp Met Lys Phe Val Ser Asp Gly Ser Ile Asn Lys
 530 535 540
 Ala Gly Phe Ala Ala Asn Phe Phe Lys Glu Val Asp Glu Cys Ser Trp
 545 550 555 560
 Pro Asp His Gly Gly Cys Glu His Arg Cys Val Asn Thr Leu Gly Ser
 565 570 575
 Tyr Lys Cys Ala Cys Asp Pro Gly Tyr Glu Leu Ala Ala Asp Lys Lys
 580 585 590
 Met Cys Glu Val Ala Cys Gly Gly Phe Ile Thr Lys Leu Asn Gly Thr
 595 600 605
 Ile Thr Ser Pro Gly Trp Pro Lys Glu Tyr Pro Thr Asn Lys Asn Cys
 610 615 620
 Val Trp Gln Val Val Ala Pro Thr Gln Tyr Arg Ile Ser Leu Gln Phe
 625 630 635 640
 Glu Val Phe Glu Leu Glu Gly Asn Asp Val Cys Lys Tyr Asp Phe Val
 645 650 655
 Glu Val Arg Ser Gly Leu Ser Pro Asp Ala Lys Leu His Gly Arg Phe
 660 665 670
 Cys Gly Ser Glu Thr Pro Glu Val Ile Thr Ser Gln Ser Asn Asn Met
 675 680 685
 Arg Val Glu Phe Lys Ser Asp Asn Thr Val Ser Lys Arg Gly Phe Arg
 690 695 700
 Ala His Phe Phe Ser Asp Lys Asp Glu Cys Ala Lys Asp Asn Gly Gly
 705 710 715 720

Cys Gln His Glu Cys Val Asn Thr Phe Gly Ser Tyr Leu Cys Arg Cys
 725 730 735
 Arg Asn Gly Tyr Trp Leu His Glu Asn Gly His Asp Cys Lys Glu Ala
 740 745 750
 Gly Cys Ala His Lys Ile Ser Ser Val Glu Gly Thr Leu Ala Ser Pro
 755 760 765
 Asn Trp Pro Asp Lys Tyr Pro Ser Arg Arg Glu Cys Thr Trp Asn Ile
 770 775 780
 Ser Ser Thr Ala Gly His Arg Val Lys Leu Thr Phe Asn Glu Phe Glu
 785 790 795 800
 Ile Glu Gln His Gln Glu Cys Ala Tyr Asp His Leu Glu Met Tyr Asp
 805 810 815
 Gly Pro Asp Ser Leu Ala Pro Ile Leu Gly Arg Phe Cys Gly Ser Lys
 820 825 830
 Lys Pro Asp Pro Thr Val Ala Ser Gly Ser Lys Cys Gly Gly Arg Leu
 835 840 845
 Lys Ala Glu Val Gln Thr Lys Glu Leu Tyr Ser His Ala Gln Phe Gly
 850 855 860
 Asp Asn Asn Tyr Pro Ser Glu Ala Arg Cys Asp Trp Val Ile Val Ala
 865 870 875 880
 Glu Asp Gly Tyr Gly Val Glu Leu Thr Phe Arg Thr Phe Glu Val Glu
 885 890 895
 Glu Glu Ala Asp Cys Gly Tyr Asp Tyr Met Glu Ala Tyr Asp Gly Tyr
 900 905 910
 Asp Ser Ser Ala Pro Arg Leu Gly Arg Phe Cys Gly Ser Gly Pro Leu
 915 920 925
 Glu Glu Ile Tyr Ser Ala Gly Asp Ser Leu Met Ile Arg Phe Arg Thr
 930 935 940
 Asp Asp Thr Ile Asn Lys Lys Gly Phe His Ala Arg Tyr Thr Ser Thr
 945 950 955 960
 Lys Phe Gln Asp Ala Leu His Met Lys Lys
 965 970

<210> 43
 <211> 1604
 <212> DNA
 <213> Homo sapiens

<400> 43
 tagattatct ctcaaacaca atttgttgc ttgcttccag gagatattga tcaacaagag 60

atgattccaa	gttgaagagg	gttggatgt	caaaaactgg	agaaaaattt	120
gttgaagagg	cctgttaggct	aataatggaa	gagggtggtt	tgaaagctac	180
gagaaggat	gtgaatggag	gcctccctgaa	caactgaaac	agcttcttga	240
agagactcg	gcgagccacc	ccataaacta	tttggactct	gtcgggatgt	300
agtgtcaaaa	cagaccaccc	aagattttc	aaccaattgt	atgctggact	360
tccttggtgg	cccgatttat	gaccgaagca	ttgaatccaa	tgatttatac	420
tccccagtgt	ttctgttagt	ggaagaagcg	gttctgaaga	aatgattga	480
tggaaagaag	gggatggaat	attnaaccct	ggggctca	tgtccatat	540
aatttagcta	gatacaaata	ttgtccctgat	attaaggaaa	aggggctgtc	600
agattaatcc	tttcacatc	tgcagagtgt	cattactcta	tgaagaaggc	660
cttgggattt	gcactgagaa	tgttgcttt	gtggaaacag	atagaggtaa	720
gaggaactgg	agaagcaagt	ctggcaagcc	agaaaagagg	gggcagcacc	780
tgtgccactt	ctggtaacaac	tgtgttggg	gcttttgacc	ctctggatga	840
atctgcgaga	ggcacagcct	ctggcttcat	gtagatgctt	cttgggggtgg	900
atgtcgagga	agcacccgaa	gcttctgcat	ggcatccaca	gggctgactc	960
aacccacaca	agatgctgat	ggctgggatc	cagtgctgt	ctctccttgt	1020
tctgacttag	aaaagagatg	ccaagagttt	gtgcctgcct	atctctggca	1080
ttttataatg	ttgttttca	aaaaatggt	acaaaattta	cccatgaaac	1140
aggaattgca	gaagcctgtg	gttcacctgg	aaagccagg	gtggtgagg	1200
ttgaggtgcc	ccatgcttagg	tgatgggagg	taccttagtag	atgaaatcaa	1260
ggattcaagt	tactgatgga	acctgaatat	gccaatattt	gcttttggta	1320
agcctcagag	agatggaaga	aggacccgag	ttctgggcaa	aacttacaca	1380
gccatttaagg	agaggatgat	gaagaaggga	agcttgatgc	tgggcttacca	1440
acaaagggtca	acttcttccg	ccaggtgggt	atcagccctc	aagtgagccg	1500
gacttctcc	tggatgagat	agacttactg	ggtaaaagaca	tgtagctgt	1560
cccaqaqqca	tagatcctat	cctgggagag	tttagatcca	gaac	1604

<210> 44
<211> 494
<212> PRT
<213> *Homo sapiens*

<400> 44
Met Ile Pro Ser Lys Lys Gly Val Val Leu Asn Gly Asp Ala Lys Ala
1 5 10 15
Gly Glu Lys Phe Val Glu Glu Ala Cys Arg Leu Ile Met Glu Glu Val
20 25 30

Val Leu Lys Ala Thr Asp Val Asn Glu Lys Val Cys Glu Trp Arg Pro
 35 40 45

Pro Glu Gln Leu Lys Gln Leu Leu Asp Leu Glu Met Arg Asp Ser Gly
 50 55 60

Glu Pro Pro His Lys Leu Leu Glu Leu Cys Arg Asp Val Ile His Tyr
 65 70 75 80

Ser Val Lys Thr Asp His Pro Arg Phe Phe Asn Gln Leu Tyr Ala Gly
85 90 95

Leu Asp Tyr Tyr Ser Leu Val Ala Arg Phe Met Thr Glu Ala Leu Asn
 100 105 110

Pro Ser Ser Tyr Thr Tyr Glu Val Ser Pro Val Phe Leu Leu Val Glu
115 120 125

Glu Ala Val Leu Lys Lys Met Ile Glu Phe Ile Gly Trp Lys Glu Gly
 130 135 140
 Asp Gly Ile Phe Asn Pro Gly Gly Ser Val Ser Asn Met Tyr Ala Met
 145 150 155 160
 Asn Leu Ala Arg Tyr Lys Tyr Cys Pro Asp Ile Lys Glu Lys Gly Leu
 165 170 175
 Ser Gly Ser Pro Arg Leu Ile Leu Phe Thr Ser Ala Glu Cys His Tyr
 180 185 190
 Ser Met Lys Lys Ala Ala Ser Phe Leu Gly Ile Gly Thr Glu Asn Val
 195 200 205
 Cys Phe Val Glu Thr Asp Arg Gly Lys Met Ile Pro Glu Glu Leu Glu
 210 215 220
 Lys Gln Val Trp Gln Ala Arg Lys Glu Gly Ala Ala Pro Phe Leu Val
 225 230 235 240
 Cys Ala Thr Ser Gly Thr Thr Val Leu Gly Ala Phe Asp Pro Leu Asp
 245 250 255
 Glu Ile Ala Asp Ile Cys Glu Arg His Ser Leu Trp Leu His Val Asp
 260 265 270
 Ala Ser Trp Gly Gly Ser Ala Leu Met Ser Arg Lys His Arg Lys Leu
 275 280 285
 Leu His Gly Ile His Arg Ala Asp Ser Val Ala Trp Asn Pro His Lys
 290 295 300
 Met Leu Met Ala Gly Ile Gln Cys Cys Ala Leu Leu Val Lys Asp Lys
 305 310 315 320
 Ser Asp Leu Glu Lys Arg Cys Gln Glu Phe Val Pro Ala Tyr Leu Trp
 325 330 335
 Gln Glu Asp Lys Phe Tyr Asn Val Ala Phe Gln Lys Asn Gly Thr Lys
 340 345 350
 Phe Thr His Glu Thr Gln Val Gly Arg Asn Cys Arg Ser Leu Trp Phe
 355 360 365
 Thr Trp Lys Ala Arg Gly Gly Glu Gly Leu Gly Trp Leu Arg Cys Pro
 370 375 380
 Met Leu Gly Asp Gly Arg Tyr Leu Val Asp Glu Ile Lys Lys Arg Glu
 385 390 395 400
 Gly Phe Lys Leu Leu Met Glu Pro Glu Tyr Ala Asn Ile Cys Phe Trp
 405 410 415
 Tyr Ile Pro Pro Ser Leu Arg Glu Met Glu Glu Gly Pro Glu Phe Trp
 420 425 430

Ala Lys Leu Thr Gln Val Ala Pro Ala Ile Lys Glu Arg Met Met Lys		
435	440	445
Lys Gly Ser Leu Met Leu Gly Tyr Gln Pro His Phe Thr Lys Val Asn		
450	455	460
Phe Phe Arg Gln Val Val Ile Ser Pro Gln Val Ser Arg Glu Asp Met		
465	470	475
Asp Phe Leu Leu Asp Glu Ile Asp Leu Leu Gly Lys Asp Met		
485	490	

<210> 45
<211> 3564
<212> PRT
<213> Mus musculus

<400> 45		
Met Thr Ala Trp Arg Lys Phe Lys Ser Leu Leu Leu Pro Leu Val Leu		
1	5	10
Ala Val Leu Cys Ala Gly Leu Leu Thr Ala Ala Lys Gly Gln Asn Cys		
20	25	30
Gly Gly Leu Val Gln Gly Pro Asn Gly Thr Ile Glu Ser Pro Gly Phe		
35	40	45
Pro His Gly Tyr Pro Asn Tyr Ala Asn Cys Thr Trp Ile Ile Ile Thr		
50	55	60
Gly Glu Arg Asn Arg Ile Gln Leu Ser Phe His Thr Phe Ala Leu Glu		
65	70	75
Glu Asp Phe Asp Ile Leu Ser Val Tyr Asp Gly Gln Pro Gln Gln Gly		
85	90	95
Asn Leu Lys Val Arg Leu Ser Gly Phe Gln Leu Pro Ser Ser Ile Val		
100	105	110
Ser Thr Gly Ser Leu Leu Thr Leu Trp Phe Thr Thr Asp Phe Ala Val		
115	120	125
Ser Ala Gln Gly Phe Lys Ala Met Tyr Glu Val Leu Pro Ser His Thr		
130	135	140
Cys Gly Asn Pro Gly Glu Ile Leu Lys Gly Val Leu His Gly Thr Arg		
145	150	155
Phe Asn Ile Gly Asp Lys Ile Arg Tyr Ser Cys Leu Ser Gly Tyr Ile		
165	170	175
Leu Glu Gly His Ala Ile Leu Thr Cys Ile Val Ser Pro Gly Asn Gly		
180	185	190
Ala Ser Trp Asp Phe Pro Ala Pro Phe Cys Arg Ala Glu Gly Ala Cys		

195	200	205
Gly Gly Thr Leu Arg Gly	Thr Ser Gly Ser Ile Ser Ser Pro His Phe	
210	215	220
Pro Ser Glu Tyr Asp Asn Asn Ala Asp Cys	Thr Trp Thr Ile Leu Ala	
225	230	235
Glu Pro Gly Asp Thr Ile Ala Leu Val	Phe Thr Asp Phe Gln Leu Glu	
245	250	255
Glu Gly Tyr Asp Phe Leu Glu Ile Ser Gly	Thr Glu Ala Pro Ser Ile	
260	265	270
Trp Leu Thr Gly Met Asn Leu Pro Ser Pro Val Ile Ser Ser Lys Asn		
275	280	285
Trp Leu Arg Leu His Phe Thr Ser Asp Ser Asn His Arg Arg Lys Gly		
290	295	300
Phe Asn Ala Gln Phe Gln Val Lys Lys Ala Ile Glu Leu Lys Ser Arg		
305	310	315
Gly Val Lys Met Leu Pro Ser Lys Asp Ser Ser His Lys Asn Ser Val		
325	330	335
Leu Thr Gln Gly Gly Val Ser Leu Ile Ser Asp Met Cys Pro Asp Pro		
340	345	350
Gly Ile Pro Asp Asn Gly Arg Arg Ala Gly Ser Asp Phe Arg Val Gly		
355	360	365
Ala Asn Val Gln Phe Ser Cys Glu Asp Asn Tyr Val Leu Gln Gly Ala		
370	375	380
Lys Gly Ile Thr Cys Gln Arg Val Thr Glu Thr Leu Ala Ala Trp Asn		
385	390	395
Asp His Arg Pro Ile Cys Arg Ala Arg Thr Cys Gly Ser Asn Leu Arg		
405	410	415
Gly Pro Ser Gly Val Ile Thr Ser Pro Asn Tyr Pro Val Gln Tyr Glu		
420	425	430
Asp Asn Ala His Cys Val Trp Val Ile Thr Thr Thr Asp Pro Asp Lys		
435	440	445
Val Ile Lys Leu Ala Phe Glu Glu Phe Glu Leu Glu Arg Gly Tyr Asp		
450	455	460
Thr Leu Thr Val Gly Asp Ala Gly Lys Val Gly Asp Thr Arg Ser Val		
465	470	475
Leu Tyr Val Leu Thr Gly Ser Ser Val Pro Asp Leu Ile Val Ser Met		
485	490	495
Ser Asn Gln Met Trp Leu His Leu Gln Ser Asp Asp Ser Ile Gly Ser		

500	505	510
Pro Gly Phe Lys Ala Val Tyr Gln Glu Ile Glu Lys Gly Gly Cys Gly		
515	520	525
Asp Pro Gly Ile Pro Ala Tyr Gly Lys Arg Thr Gly Ser Ser Phe Leu		
530	535	540
His Gly Asp Thr Leu Thr Phe Glu Cys Gln Ala Ala Phe Glu Leu Val		
545	550	555
Gly Glu Arg Val Ile Thr Cys Gln Lys Asn Asn Gln Trp Ser Gly Asn		
565	570	575
Lys Pro Ser Cys Val Phe Ser Cys Phe Phe Asn Phe Thr Ala Pro Ser		
580	585	590
Gly Ile Ile Leu Ser Pro Asn Tyr Pro Glu Glu Tyr Gly Asn Asn Met		
595	600	605
Asn Cys Val Trp Leu Ile Ile Ser Glu Pro Gly Ser Arg Ile His Leu		
610	615	620
Ile Phe Lys Asp Phe Asp Val Glu Pro Gln Phe Asp Phe Leu Ala Val		
625	630	640
Lys Asp Asp Gly Ile Ser Asp Ile Thr Val Leu Gly Thr Phe Ser Gly		
645	650	655
Asn Glu Val Pro Ala Gln Leu Ala Ser Ser Gly His Ile Val Arg Leu		
660	665	670
Glu Phe Gln Ser Asp His Ser Thr Thr Gly Arg Gly Phe Asn Ile Thr		
675	680	685
Tyr Thr Thr Phe Gly Gln Asn Glu Cys His Asp Pro Gly Ile Pro Val		
690	695	700
Asn Gly Arg Arg Phe Gly Asp Arg Phe Leu Leu Gly Ser Ser Val Ser		
705	710	720
Phe His Cys Asp Asp Gly Phe Val Lys Thr Gln Gly Ser Glu Ser Ile		
725	730	735
Thr Cys Ile Leu Gln Asp Gly Asn Val Val Trp Ser Ser Thr Val Pro		
740	745	750
Arg Cys Glu Ala Pro Cys Gly Gly His Leu Thr Ala Ser Ser Gly Val		
755	760	765
Ile Leu Pro Pro Gly Trp Pro Gly Tyr Tyr Lys Asp Ser Leu Asn Cys		
770	775	780
Glu Trp Val Ile Glu Ala Lys Pro Gly His Ser Ile Lys Ile Thr Phe		
785	790	795
Asp Arg Phe Gln Thr Glu Val Asn Tyr Asp Thr Leu Glu Val Arg Asp		

805	810	815
Gly Pro Thr Ser Ser Ser Pro Leu Ile Gly Glu Tyr His Gly Thr Gln		
820	825	830
Ala Pro Gln Phe Leu Ile Ser Thr Gly Asn Tyr Met Tyr Leu Leu Phe		
835	840	845
Thr Thr Asp Ser Ser Arg Ala Ser Val Gly Phe Leu Ile His Tyr Glu		
850	855	860
Ser Val Thr Leu Glu Ser Asp Ser Cys Leu Asp Pro Gly Ile Pro Val		
865	870	875
Asn Gly Gln Arg His Gly Ser Asn Phe Gly Ile Arg Ser Thr Val Thr		
885	890	895
Phe Ser Cys Asp Pro Gly Tyr Thr Leu Ser Asp Asp Glu Pro Leu Val		
900	905	910
Cys Glu Lys Asn His Gln Trp Asn His Ala Leu Pro Ser Cys Asp Ala		
915	920	925
Leu Cys Gly Gly Tyr Ile His Gly Lys Ser Gly Thr Val Leu Ser Pro		
930	935	940
Gly Phe Pro Asp Phe Tyr Pro Asn Ser Leu Asn Cys Thr Trp Thr Ile		
945	950	955
960		
Glu Val Ser His Gly Lys Gly Val Gln Met Asn Phe His Thr Phe His		
965	970	975
Leu Glu Ser Ser His Asp Tyr Leu Leu Ile Thr Glu Asp Gly Ser Phe		
980	985	990
Ser Glu Pro Val Ala Arg Leu Thr Gly Ser Val Leu Pro His Thr Ile		
995	1000	1005
1005		
Lys Ala Gly Leu Phe Gly Asn Phe Thr Ala Gln Leu Arg Phe Ile Ser		
1010	1015	1020
Asp Phe Ser Ile Ser Tyr Glu Gly Phe Asn Ile Thr Phe Ala Glu Tyr		
1025	1030	1035
1040		
Asp Leu Glu Pro Cys Asp Asp Pro Gly Val Pro Ala Phe Ser Arg Arg		
1045	1050	1055
Ile Gly Phe Gln Phe Gly Val Gly Asp Thr Leu Ala Phe Thr Cys Phe		
1060	1065	1070
Gln Gly Tyr Arg Leu Glu Gly Ala Thr Lys Leu Thr Cys Leu Gly Gly		
1075	1080	1085
Gly Arg Arg Val Trp Ser Ala Pro Leu Pro Arg Cys Val Ala Glu Cys		
1090	1095	1100
Gly Ala Ser Val Lys Gly Asn Glu Gly Thr Leu Leu Ser Pro Asn Phe		

1105	1110	1115	1120
Pro Ser His Tyr Asp Asn Asn His Glu Cys Ile Tyr Lys Ile Glu Thr			
1125		1130	1135
Glu Ala Gly Lys Gly Ile His Leu Arg Ala Arg Thr Phe Gln Leu Phe			
1140	1145		1150
Glu Gly Asp Thr Leu Lys Val Tyr Asp Gly Lys Asp Ser Ser Ser Arg			
1155	1160	1165	
Ser Leu Gly Val Phe Thr Arg Ser Glu Phe Met Gly Leu Val Leu Asn			
1170	1175	1180	
Ser Thr Ser Asn Tyr Leu Arg Leu Glu Phe Asn Thr Asn Gly Ser Asp			
1185	1190	1195	1200
Thr Ala Gln Gly Phe Gln Leu Thr Tyr Thr Ser Phe Asp Leu Val Lys			
1205	1210		1215
Cys Glu Asp Pro Gly Ile Pro Asn Tyr Gly Tyr Arg Ile Arg Asp Asp			
1220	1225	1230	
Gly His Phe Thr Asp Thr Val Val Leu Tyr Ser Cys Asn Pro Gly Tyr			
1235	1240	1245	
Ala Met His Gly Ser Ser Thr Leu Thr Cys Leu Ser Gly Asp Arg Arg			
1250	1255	1260	
Val Trp Asp Lys Pro Met Pro Ser Cys Val Ala Glu Cys Gly Gly Leu			
1265	1270	1275	1280
Val His Ala Ala Thr Ser Gly Arg Ile Leu Ser Pro Gly Tyr Pro Ala			
1285	1290		1295
Pro Tyr Asp Asn Asn Leu His Cys Thr Trp Thr Ile Glu Ala Asp Pro			
1300	1305	1310	
Gly Lys Thr Ile Ser Leu His Phe Ile Val Phe Asp Thr Glu Thr Ala			
1315	1320	1325	
His Asp Ile Leu Lys Val Trp Asp Gly Pro Val Asp Ser Asn Ile Leu			
1330	1335	1340	
Leu Lys Glu Trp Ser Gly Ser Ala Leu Pro Glu Asp Ile His Ser Thr			
1345	1350	1355	1360
Phe Asn Ser Leu Thr Leu Gln Phe Asp Ser Asp Phe Phe Ile Ser Lys			
1365	1370	1375	
Ser Gly Phe Ser Ile Gln Phe Ser Thr Ser Ile Ala Ser Thr Cys Asn			
1380	1385	1390	
Asp Pro Gly Met Pro Gln Asn Gly Thr Arg Tyr Gly Asp Ser Arg Glu			
1395	1400	1405	
Pro Gly Asp Thr Ile Thr Phe Gln Cys Asp Pro Gly Tyr Gln Leu Gln			

1410	1415	1420
Gly Pro Ala Lys Ile Thr Cys Val Gln Leu Asn Asn Arg Phe Phe Trp		
1425	1430	1435
Gln Pro Asp Pro Pro Ser Cys Ile Ala Ala Cys Gly Gly Asn Leu Thr		
1445	1450	1455
Gly Pro Ala Gly Val Ile Leu Ser Pro Asn Tyr Pro Gln Pro Tyr Pro		
1460	1465	1470
Pro Gly Lys Glu Cys Asp Trp Arg Ile Lys Val Asn Pro Asp Phe Val		
1475	1480	1485
Ile Ala Leu Ile Phe Lys Ser Phe Ser Met Glu Pro Ser Tyr Asp Phe		
1490	1495	1500
Leu His Ile Tyr Glu Gly Glu Asp Ser Asn Ser Pro Leu Ile Gly Ser		
1505	1510	1515
Phe Gln Gly Ser Gln Ala Pro Glu Arg Ile Glu Ser Ser Gly Asn Ser		
1525	1530	1535
Leu Phe Leu Ala Phe Arg Ser Asp Ala Ser Val Gly Leu Ser Gly Phe		
1540	1545	1550
Ala Ile Glu Phe Lys Glu Lys Pro Arg Glu Ala Cys Phe Asp Pro Gly		
1555	1560	1565
Asn Ile Met Asn Gly Thr Arg Ile Gly Thr Asp Phe Lys Leu Gly Ser		
1570	1575	1580
Thr Val Thr Tyr Gln Cys Asp Ser Gly Tyr Lys Ile Val Asp Pro Ser		
1585	1590	1595
Ser Ile Glu Cys Val Thr Gly Ala Asp Gly Lys Pro Ser Trp Asp Arg		
1605	1610	1615
Ala Leu Pro Ala Cys Gln Ala Pro Cys Gly Gly Gln Tyr Thr Gly Ser		
1620	1625	1630
Glu Gly Val Val Leu Ser Pro Asn Tyr Pro His Asn Tyr Thr Ala Gly		
1635	1640	1645
Gln Met Cys Val Tyr Ser Ile Thr Val Pro Lys Glu Phe Val Val Phe		
1650	1655	1660
Gly Gln Phe Ala Tyr Phe Gln Thr Ala Leu Asn Asp Leu Ala Glu Leu		
1665	1670	1675
Phe Asp Gly Thr His Pro Gln Ala Arg Leu Leu Ser Ser Leu Ser Gly		
1685	1690	1695
Ser His Ser Gly Glu Thr Leu Pro Leu Ala Thr Ser Asn Gln Ile Leu		
1700	1705	1710
Leu Arg Phe Ser Ala Lys Ser Gly Ala Ser Ala Arg Gly Phe His Phe		

1715	1720	1725	
Val Tyr Gln Ala Val Pro Arg Thr Ser Asp Thr Gln Cys Ser Ser Val			
1730	1735	1740	
Pro Glu Pro Arg Tyr Gly Arg Arg Ile Gly Ser Glu Phe Ser Ala Gly			
1745	1750	1760	
Ser Ile Val Arg Phe Glu Cys Asn Pro Gly Tyr Leu Leu Gln Gly Ser			
1765	1770	1775	
Thr Ala Ile Arg Cys Gln Ser Val Pro Asn Ala Leu Ala Gln Trp Asn			
1780	1785	1790	
Asp Thr Ile Pro Ser Cys Val Val Pro Cys Ser Gly Asn Phe Thr Gln			
1795	1800	1805	
Arg Arg Gly Thr Ile Leu Ser Pro Gly Tyr Pro Glu Pro Tyr Gly Asn			
1810	1815	1820	
Asn Leu Asn Cys Val Trp Lys Ile Ile Val Ser Glu Gly Ser Gly Ile			
1825	1830	1835	1840
Gln Ile Gln Val Ile Ser Phe Ala Thr Glu Gln Asn Trp Asp Ser Leu			
1845	1850	1855	
Glu Ile His Asp Gly Gly Asp Met Thr Ala Pro Arg Leu Gly Ser Phe			
1860	1865	1870	
Ser Gly Thr Thr Val Pro Ala Leu Leu Asn Ser Thr Ser Asn Gln Leu			
1875	1880	1885	
Cys Leu His Phe Gln Ser Asp Ile Ser Val Ala Ala Gly Phe His			
1890	1895	1900	
Leu Glu Tyr Lys Thr Val Gly Leu Ala Ala Cys Gln Glu Pro Ala Leu			
1905	1910	1915	1920
Pro Ser Asn Gly Ile Lys Ile Gly Asp Arg Tyr Met Val Asn Asp Val			
1925	1930	1935	
Leu Ser Phe Gln Cys Glu Pro Gly Tyr Thr Leu Gln Gly Arg Ser His			
1940	1945	1950	
Ile Ser Cys Met Pro Gly Thr Val Arg Arg Trp Asn Tyr Pro Ser Pro			
1955	1960	1965	
Leu Cys Ile Ala Thr Cys Gly Gly Thr Leu Thr Ser Met Ser Gly Val			
1970	1975	1980	
Ile Leu Ser Pro Gly Phe Pro Gly Ser Tyr Pro Asn Asn Leu Asp Cys			
1985	1990	1995	2000
Thr Trp Lys Ile Ser Leu Pro Ile Gly Tyr Gly Ala His Ile Gln Phe			
2005	2010	2015	
Leu Asn Phe Ser Thr Glu Ala Asn His Asp Tyr Leu Glu Ile Gln Asn			

2020	2025	2030
Gly Pro Tyr His Ser Ser Pro Met Met Gly Gln Phe Ser Gly Pro Asp		
2035	2040	2045
Leu Pro Thr Ser Leu Leu Ser Thr Thr His Glu Thr Leu Ile Arg Phe		
2050	2055	2060
Tyr Ser Asp His Ser Gln Asn Arg Gln Gly Phe Lys Leu Ser Tyr Gln		
2065	2070	2075
Ala Tyr Glu Leu Gln Asn Cys Pro Asp Pro Pro Ala Phe Gln Asn Gly		
2085	2090	2095
Phe Met Ile Asn Ser Asp Tyr Ser Val Gly Gln Ser Ile Ser Phe Glu		
2100	2105	2110
Cys Tyr Pro Gly Tyr Ile Leu Leu Gly His Pro Val Leu Thr Cys Gln		
2115	2120	2125
His Gly Thr Asp Arg Asn Trp Asn Tyr Pro Phe Pro Arg Cys Asp Ala		
2130	2135	2140
Pro Cys Gly Tyr Asn Val Thr Ser Gln Asn Gly Thr Ile Tyr Ser Pro		
2145	2150	2155
Gly Phe Pro Asp Glu Tyr Pro Ile Leu Lys Asp Cys Leu Trp Leu Val		
2165	2170	2175
Thr Val Pro Pro Gly His Gly Val Tyr Ile Asn Phe Thr Leu Leu Gln		
2180	2185	2190
Thr Glu Ala Val Asn Asp Tyr Ile Ala Val Trp Asp Gly Pro Asp Gln		
2195	2200	2205
Asn Ser Pro Gln Leu Gly Val Phe Ser Gly Asn Thr Ala Pro Glu Thr		
2210	2215	2220
Ala Tyr Ser Ser Thr Asn Gln Val Leu Leu Lys Phe His Ser Asp Phe		
2225	2230	2235
Ser Asn Gly Gly Phe Phe Val Leu Asn Phe His Ala Phe Gln Leu Lys		
2245	2250	2255
Arg Cys Pro Pro Pro Ala Val Pro Gln Ala Asp Leu Leu Thr Glu		
2260	2265	2270
Asp Glu Asp Phe Glu Ile Gly Asp Phe Val Lys Tyr Gln Cys His Pro		
2275	2280	2285
Gly Tyr Thr Leu Leu Gly Ser Asp Thr Leu Thr Cys Lys Leu Ser Ser		
2290	2295	2300
Gln Leu Leu Phe Gln Gly Ser Pro Pro Thr Cys Glu Ala Gln Cys Pro		
2305	2310	2315
Ala Asn Glu Val Arg Thr Glu Ser Ser Gly Val Ile Leu Ser Pro Gly		

2325	2330	2335
Tyr Pro Gly Asn Tyr Phe Asn Ser Gln Thr Cys Ala Trp Ser Ile Lys		
2340	2345	2350
Val Lys Pro Asn Phe Asn Ile Thr Leu Phe Val Asp Thr Phe Gln Ser		
2355	2360	2365
Glu Lys Gln Phe Asp Ala Leu Glu Val Phe Asp Gly Ser Ser Gly Arg		
2370	2375	2380
Ser Pro Leu Leu Val Val Leu Ser Gly Asn His Thr Glu Gln Ser Asn		
2385	2390	2395
Phe Thr Ser Arg Ser Asn His Leu Tyr Leu Arg Trp Ser Thr Asp His		
2405	2410	2415
Ala Thr Ser Lys Lys Gly Phe Lys Ile Arg Tyr Ala Ala Pro Tyr Cys		
2420	2425	2430
Ser Leu Thr Ser Thr Leu Arg Asn Gly Gly Ile Leu Asn Lys Thr Ala		
2435	2440	2445
Gly Ala Val Gly Ser Lys Val His Tyr Phe Cys Lys Pro Gly Tyr Arg		
2450	2455	2460
Met Ile Gly His Ser Asn Ala Thr Cys Arg Arg Asn Pro Val Gly Val		
2465	2470	2475
Tyr Gln Trp Asp Ser Met Ala Pro Leu Cys Gln Ala Val Ser Cys Gly		
2485	2490	2495
Ile Pro Glu Ala Pro Gly Asn Gly Ser Phe Thr Gly Asn Glu Phe Thr		
2500	2505	2510
Leu Asp Ser Lys Val Thr Tyr Glu Cys Asn Glu Gly Phe Lys Leu Asp		
2515	2520	2525
Ala Ser Gln Glu Ala Thr Thr Val Cys Gln Glu Asp Gly Leu Trp Ser		
2530	2535	2540
Asn Arg Gly Lys Pro Pro Thr Cys Lys Pro Val Pro Cys Pro Ser Ile		
2545	2550	2555
Glu Gly Gln Leu Ser Glu His Val Leu Trp Arg Leu Val Ser Gly Ser		
2565	2570	2575
Leu Asn Glu Tyr Gly Ala Gln Val Leu Leu Ser Cys Ser Pro Gly Tyr		
2580	2585	2590
Phe Leu Gln Gly Gln Arg Leu Leu Gln Cys Gln Ala Asn Gly Thr Trp		
2595	2600	2605
Ser Thr Glu Glu Asp Arg Pro Arg Cys Lys Val Ile Ser Cys Gly Ser		
2610	2615	2620
Leu Ser Phe Pro Pro Asn Gly Asn Lys Ile Gly Thr Leu Thr Ile Tyr		

2625	2630	2635	2640
Gly Ala Thr Ala Ile Phe Thr Cys Asn Thr Gly Tyr Thr Leu Val Gly			
2645	2650		2655
Ser His Val Arg Glu Cys Leu Ala Asn Gly Leu Trp Ser Gly Ser Glu			
2660	2665		2670
Thr Arg Cys Leu Ala Gly His Cys Gly Ser Pro Asp Pro Ile Val Asn			
2675	2680		2685
Gly His Ile Ser Gly Asp Gly Phe Ser Tyr Arg Asp Thr Val Val Tyr			
2690	2695		2700
Gln Cys Asn Pro Gly Phe Arg Leu Val Gly Thr Ser Val Arg Ile Cys			
2705	2710	2715	2720
Cys Arg Thr Thr Ser Gly Arg Gly Arg Leu Thr Val Cys Val Pro Ile			
2725	2730		2735
Thr Cys Gly His Pro Gly Asn Pro Ala His Gly Leu Thr Asn Gly Thr			
2740	2745		2750
Glu Phe Asn Leu Asn Asp Leu Val Asn Phe Thr Cys His Thr Gly Tyr			
2755	2760		2765
Arg Leu Gln Gly Ala Ser Arg Ala Gln Cys Arg Ser Asn Gly Gln Trp			
2770	2775		2780
Ser Ser Pro Leu Pro Ile Cys Arg Val Val Asn Cys Ser Asp Pro Gly			
2785	2790	2795	2800
Ser Val Glu Asn Ala Val Arg His Gly Gln Gln Asn Phe Pro Glu Ser			
2805	2810		2815
Phe Glu Tyr Gly Thr Ser Val Met Tyr His Cys Lys Thr Gly Phe Tyr			
2820	2825		2830
Leu Leu Gly Ser Ser Ala Leu Thr Cys Met Ala Ser Gly Leu Trp Asp			
2835	2840		2845
Arg Ser Leu Pro Lys Cys Leu Ala Ile Ser Cys Gly His Pro Gly Val			
2850	2855		2860
Pro Ala Asn Ala Val Leu Thr Gly Glu Leu Phe Thr Tyr Gly Ala Thr			
2865	2870	2875	2880
Val Gln Tyr Ser Cys Lys Gly Gln Ile Leu Thr Gly Asn Ser Thr			
2885	2890		2895
Arg Val Cys Gln Glu Asp Ser His Trp Ser Gly Ser Leu Pro His Cys			
2900	2905		2910
Ser Gly Asn Ser Pro Gly Phe Cys Gly Asp Pro Gly Thr Pro Ala His			
2915	2920		2925
Gly Ser Arg Leu Gly Asp Glu Phe Lys Thr Lys Ser Leu Leu Arg Phe			

2930	2935	2940
Ser Cys Glu Met Gly His Gln Leu Arg Gly Phe Ala Glu Arg Thr Cys		
2945	2950	2955
2960		
Leu Val Asn Gly Ser Trp Ser Gly Val Gln Pro Val Cys Glu Ala Val		
2965	2970	2975
Ser Cys Gly Asn Pro Gly Thr Pro Thr Asn Gly Met Ile Leu Ser Ser		
2980	2985	2990
Asp Gly Ile Leu Phe Ser Ser Val Ile Tyr Ala Cys Trp Glu Gly		
2995	3000	3005
Tyr Lys Thr Ser Gly Leu Met Thr Arg His Cys Thr Ala Asn Gly Thr		
3010	3015	3020
Trp Thr Gly Thr Ala Pro Asp Cys Thr Ile Ile Ser Cys Gly Asp Pro		
3025	3030	3035
3040		
Gly Thr Leu Pro Asn Gly Ile Gln Phe Gly Thr Asp Phe Thr Phe Asn		
3045	3050	3055
Lys Thr Val Ser Tyr Gln Cys Asn Pro Gly Tyr Leu Met Glu Pro Pro		
3060	3065	3070
Thr Ser Pro Thr Ile Arg Cys Thr Lys Asp Gly Thr Trp Asn Gln Ser		
3075	3080	3085
Arg Pro Leu Cys Lys Ala Val Leu Cys Asn Gln Pro Pro Pro Val Pro		
3090	3095	3100
Asn Gly Lys Val Glu Gly Ser Asp Phe Arg Trp Gly Ala Ser Ile Ser		
3105	3110	3115
3120		
Tyr Ser Cys Val Asp Gly Tyr Gln Leu Ser His Ser Ala Ile Leu Ser		
3125	3130	3135
Cys Glu Gly Arg Gly Val Trp Lys Gly Glu Val Pro Gln Cys Leu Pro		
3140	3145	3150
Val Phe Cys Gly Asp Pro Gly Thr Pro Ala Glu Gly Arg Leu Ser Gly		
3155	3160	3165
Lys Ser Phe Thr Phe Lys Ser Glu Val Phe Ile Gln Cys Lys Pro Pro		
3170	3175	3180
Phe Val Leu Val Gly Ser Ser Arg Arg Thr Cys Gln Ala Asp Gly Ile		
3185	3190	3195
3200		
Trp Ser Gly Ile Gln Pro Thr Cys Ile Asp Pro Ala His Thr Ala Cys		
3205	3210	3215
Pro Asp Pro Gly Thr Pro His Phe Gly Ile Gln Asn Ser Ser Lys Gly		
3220	3225	3230
Tyr Glu Val Gly Ser Thr Val Phe Phe Arg Cys Arg Lys Gly Tyr His		

3235	3240	3245
Ile Gln Gly Ser Thr Thr Arg Thr Cys Leu Ala Asn Leu Thr Trp Ser		
3250	3255	3260
Gly Ile Gln Thr Glu Cys Ile Pro His Ala Cys Arg Gln Pro Glu Thr		
3265	3270	3275
3280		
Pro Ala His Ala Asp Val Arg Ala Ile Asp Leu Pro Ala Phe Gly Tyr		
3285	3290	3295
Thr Leu Val Tyr Thr Cys His Pro Gly Phe Phe Leu Ala Gly Gly Ser		
3300	3305	3310
Glu His Arg Thr Cys Lys Ala Asp Met Lys Trp Thr Gly Lys Ser Pro		
3315	3320	3325
Val Cys Lys Ser Lys Gly Val Arg Glu Val Asn Glu Thr Val Thr Lys		
3330	3335	3340
Thr Pro Val Pro Ser Asp Val Phe Phe Ile Asn Ser Val Trp Lys Gly		
3345	3350	3355
3360		
Tyr Tyr Glu Tyr Leu Gly Lys Arg Gln Pro Ala Thr Leu Thr Val Asp		
3365	3370	3375
Trp Phe Asn Ala Thr Ser Ser Lys Val Asn Ala Thr Phe Thr Ala Ala		
3380	3385	3390
Ser Arg Val Gln Leu Glu Leu Thr Gly Val Tyr Lys Lys Glu Glu Ala		
3395	3400	3405
His Leu Leu Leu Lys Ala Phe His Ile Lys Gly Pro Ala Asp Ile Phe		
3410	3415	3420
Val Ser Lys Phe Glu Asn Asp Asn Trp Gly Leu Asp Gly Tyr Val Ser		
3425	3430	3435
3440		
Ser Gly Leu Glu Arg Gly Phe Ser Phe Gln Gly Asp Ile His Gly		
3445	3450	3455
Lys Asp Phe Gly Lys Phe Lys Leu Glu Arg Gln Asp Pro Ser Asn Ser		
3460	3465	3470
Asp Ala Asp Ser Ser Asn His Tyr Gln Gly Thr Ser Ser Gly Ser Val		
3475	3480	3485
Ala Ala Ala Ile Leu Val Pro Phe Phe Ala Leu Ile Leu Ser Gly Phe		
3490	3495	3500
Ala Phe Tyr Leu Tyr Lys His Arg Thr Arg Pro Lys Val Gln Tyr Asn		
3505	3510	3515
3520		
Gly Tyr Ala Gly His Glu Asn Ser Asn Gly Gln Ala Ser Phe Glu Asn		
3525	3530	3535
Pro Met Tyr Asp Thr Asn Leu Lys Pro Thr Glu Ala Lys Ala Val Arg		

3540

3545

3550

Phe Asp Thr Thr Leu Asn Thr Val Cys Thr Val Val
3555 3560

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<212> PRT
<213> *Homo sapiens*

<400> 46
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Leu Val Leu Cys Ala Arg Leu Leu Thr Ala Ala Lys Gly Gln Asn Cys
20 25 30

Gly Gly Leu Val Gln Gly Pro Asn Gly Thr Ile Glu Ser Pro Gly Phe
35 40 45

Pro His Gly Tyr Pro Asn Tyr Ala Asn Cys Thr Trp Ile Ile Ile Thr
 50 55 60

Glu Asn Phe Asp Ile Leu Ser Val Tyr Asp Gly Gln Pro Gln Gln Gly
 85 90 95

Asn	Leu	Lys	Val	Arg	Leu	Ser	Gly	Phe	Gln	Leu	Pro	Ser	Ser	Ile	Val
			100					105						110	

Ser Thr Gly Ser Ile Leu Thr Leu Trp Phe Thr Thr Asp Phe Ala Val
 115 120 125

Ser Ala Gln Gly Phe Lys Ala Leu Tyr Glu Val Leu Pro Ser His Thr
 130 135 140

Cys Gly Asn Pro Gly Glu Ile Leu Lys Gly Val Leu His Gly Thr Arg
145 150 155 160

Phe Asn Ile Gly Asp Lys Ile Arg Tyr Ser Cys Leu Pro Gly Tyr Ile
165 170 175

Leu Glu Gly His Ala Ile Leu Thr Cys Ile Val Ser Pro Gly Asn Gly
180 185 190

Ala Ser Trp Asp Phe Pro Ala Pro Phe Cys Arg Ala Glu Gly Ala Cys
 195 200 205

Gly Gly Thr Leu Arg Gly Thr Ser Ser Ser Ile Ser Ser Pro His Phe
212 215 220

Pro Ser Glu Tyr Glu Asn Asn Ala Asp Cys Thr Trp Thr Ile Leu Ala
 225 230 235 240

Glu Pro Gly Asp Thr Ile Ala Leu Val Phe Thr Asp Phe Gln Leu Glu
 245 250 255
 Glu Gly Tyr Asp Phe Leu Glu Ile Ser Gly Thr Glu Ala Pro Ser Ile
 260 265 270
 Trp Leu Thr Gly Met Asn Leu Pro Ser Pro Val Ile Ser Ser Lys Asn
 275 280 285
 Trp Leu Arg Leu His Phe Thr Ser Asp Ser Asn His Arg Arg Lys Gly
 290 295 300
 Phe Asn Ala Gln Phe Gln Val Lys Lys Ala Ile Glu Leu Lys Ser Arg
 305 310 315 320
 Gly Val Lys Met Leu Pro Ser Lys Asp Gly Ser His Lys Asn Ser Val
 325 330 335
 Leu Ser Gln Gly Gly Val Ala Leu Val Ser His Met Cys Leu Asp Pro
 340 345 350
 Gly Ile Pro Glu Asn Gly Arg Arg Ala Gly Ser Asp Phe Ser Arg Val
 355 360 365
 Gly Ala Asn Val Gln Phe Ser Cys Glu Asp Asn Tyr Val Leu Gln Gly
 370 375 380
 Ser Lys Ser Ile Thr Cys Gln Arg Val Thr Glu Thr Leu Ala Ala Trp
 385 390 395 400
 Ser Asp His Arg Pro Ile Cys Arg Ala Arg Thr Cys Gly Ser Asn Leu
 405 410 415
 Arg Gly Pro Ser Gly Val Ile Thr Ser Pro Asn Tyr Pro Val Gln Tyr
 420 425 430
 Glu Asp Asn Ala His Cys Val Trp Val Ile Thr Thr Thr Asp Pro Asp
 435 440 445
 Lys Val Ile Lys Leu Ala Phe Glu Glu Phe Glu Leu Glu Arg Gly Tyr
 450 455 460
 Asp Thr Leu Thr Val Gly Asp Ala Gly Lys Val Gly Asp Thr Arg Ser
 465 470 475 480
 Val Leu Tyr Val Leu Thr Gly Ser Ser Val Pro Asp Leu Ile Val Ser
 485 490 495
 Met Ser Asn Gln Met Trp Leu His Leu Gln Ser Asp Asp Ser Ile Gly
 500 505 510
 Ser Pro Gly Phe Lys Ala Val Tyr Gln Glu Ile Glu Lys Gly Gly Cys
 515 520 525
 Gly Asp Pro Gly Ile Pro Ala Tyr Gly Lys Arg Thr Gly Ser Ser Phe
 530 535 540

Leu His Gly Asp Thr Leu Thr Phe Glu Cys Pro Ala Ala Phe Glu Leu
 545 550 555 560
 Val Gly Glu Arg Val Ile Thr Cys Gln Gln Asn Asn Gln Trp Ser Gly
 565 570 575
 Asn Lys Pro Ser Cys Val Phe Ser Cys Phe Phe Asn Phe Thr Ala Ser
 580 585 590
 Ser Gly Ile Ile Leu Ser Pro Asn Tyr Pro Glu Glu Tyr Gly Asn Asn
 595 600 605
 Met Asn Cys Val Trp Leu Ile Ile Ser Glu Pro Gly Ser Arg Ile His
 610 615 620
 Leu Ile Phe Asn Asp Phe Asp Val Glu Pro Gln Phe Asp Phe Leu Ala
 625 630 635 640
 Val Lys Asp Asn Gly Ile Ser Asp Ile Thr Val Leu Gly Thr Phe Ser
 645 650 655
 Gly Asn Glu Val Pro Ser Gln Leu Ala Ser Ser Gly His Ile Val Arg
 660 665 670
 Leu Glu Phe Gln Ser Asp His Ser Thr Thr Gly Arg Gly Phe Asn Ile
 675 680 685
 Thr Tyr Thr Phe Gly Gln Asn Glu Cys His Asp Pro Gly Ile Pro
 690 695 700
 Ile Asn Gly Arg Arg Phe Gly Asp Arg Phe Leu Leu Gly Ser Ser Val
 705 710 715 720
 Ser Phe His Cys Asp Asp Gly Phe Val Lys Thr Gln Gly Ser Glu Ser
 725 730 735
 Ile Thr Cys Ile Leu Gln Asp Gly Asn Val Val Trp Ser Ser Thr Val
 740 745 750
 Pro Arg Cys Glu Ala Pro Cys Gly Gly His Leu Thr Ala Ser Ser Gly
 755 760 765
 Val Ile Leu Pro Pro Gly Trp Pro Gly Tyr Tyr Lys Asp Ser Leu His
 770 775 780
 Cys Glu Trp Ile Ile Glu Ala Lys Pro Gly His Ser Ile Lys Met Thr
 785 790 795 800
 Phe Asp Arg Phe Gln Thr Glu Val Asn Tyr Asp Thr Leu Glu Val Arg
 805 810 815
 Asp Gly Pro Ala Ser Ser Ser Pro Leu Ile Gly Glu Tyr His Gly Thr
 820 825 830
 Gln Ala Pro Gln Phe Leu Ile Ser Thr Gly Asn Phe Met Tyr Leu Leu
 835 840 845

Phe Thr Thr Asp Asn Ser Arg Ser Ser Ile Gly Phe Leu Ile His Tyr
 850 855 860
 Glu Ser Val Thr Leu Glu Ser Asp Ser Cys Leu Asp Pro Gly Ile Pro
 865 870 875 880
 Val Asn Gly His Arg His Gly Gly Asp Phe Gly Ile Arg Ser Thr Val
 885 890 895
 Thr Phe Ser Cys Asp Pro Gly Tyr Thr Leu Ser Asp Asp Glu Pro Leu
 900 905 910
 Val Cys Glu Arg Asn His Gln Trp Asn His Ala Leu Pro Ser Cys Asp
 915 920 925
 Ala Leu Cys Gly Gly Tyr Ile Gln Gly Lys Ser Gly Thr Val Leu Ser
 930 935 940
 Pro Gly Phe Pro Asp Phe Tyr Pro Asn Ser Leu Asn Cys Thr Trp Thr
 945 950 955 960
 Ile Glu Val Ser His Gly Lys Gly Val Gln Met Ile Phe His Thr Phe
 965 970 975
 His Leu Glu Ser Ser His Asp Tyr Leu Leu Ile Thr Glu Asp Gly Ser
 980 985 990
 Phe Ser Glu Pro Val Ala Arg Leu Thr Gly Ser Val Leu Pro His Thr
 995 1000 1005
 Ile Lys Ala Gly Leu Phe Gly Asn Phe Thr Ala Gln Leu Arg Phe Ile
 1010 1015 1020
 Ser Asp Phe Ser Ile Ser Tyr Glu Gly Phe Asn Ile Thr Phe Ser Glu
 1025 1030 1035 1040
 Tyr Asp Leu Glu Pro Cys Asp Asp Pro Gly Val Pro Ala Phe Ser Arg
 1045 1050 1055
 Arg Ile Gly Phe His Phe Gly Val Gly Asp Ser Leu Thr Phe Ser Cys
 1060 1065 1070
 Phe Leu Gly Tyr Arg Leu Glu Gly Ala Thr Lys Leu Thr Cys Leu Gly
 1075 1080 1085
 Gly Gly Arg Arg Val Trp Ser Ala Pro Leu Pro Arg Cys Val Ala Glu
 1090 1095 1100
 Cys Gly Ala Ser Val Lys Gly Asn Glu Gly Thr Leu Leu Ser Pro Asn
 1105 1110 1115 1120
 Phe Pro Ser Asn Tyr Asp Asn Thr His Glu Cys Ile Tyr Lys Ile Glu
 1125 1130 1135
 Thr Glu Ala Gly Lys Gly Ile His Leu Arg Thr Arg Ser Phe Gln Leu
 1140 1145 1150

Phe Glu Gly Asp Thr Leu Lys Val Tyr Asp Gly Lys Asp Ser Ser Ser
 1155 1160 1165
 Arg Pro Leu Gly Thr Phe Thr Lys Asn Glu Leu Leu Gly Leu Ile Leu
 1170 1175 1180
 Asn Ser Thr Ser Asn His Leu Trp Leu Glu Phe Asn Thr Asn Gly Ser
 1185 1190 1195 1200
 Asp Thr Asp Gln Gly Phe Gln Leu Thr Tyr Thr Ser Phe Asp Leu Val
 1205 1210 1215
 Lys Cys Glu Asp Pro Gly Ile Pro Asn Tyr Gly Tyr Arg Ile Arg Asp
 1220 1225 1230
 Glu Gly His Phe Thr Asp Thr Val Val Leu Tyr Ser Cys Asn Pro Gly
 1235 1240 1245
 Tyr Ala Met His Gly Ser Asn Thr Leu Thr Cys Leu Ser Gly Asp Arg
 1250 1255 1260
 Arg Val Trp Asp Lys Pro Leu Pro Ser Cys Ile Ala Glu Cys Gly Gly
 1265 1270 1275 1280
 Gln Ile His Ala Ala Thr Ser Gly Arg Ile Leu Ser Pro Gly Tyr Pro
 1285 1290 1295
 Ala Pro Tyr Asp Asn Asn Leu His Cys Thr Trp Ile Ile Glu Ala Asp
 1300 1305 1310
 Pro Gly Lys Thr Ile Ser Leu His Phe Ile Val Phe Asp Thr Glu Met
 1315 1320 1325
 Ala His Asp Ile Leu Lys Val Trp Asp Gly Pro Val Asp Ser Asp Ile
 1330 1335 1340
 Leu Leu Lys Glu Trp Ser Gly Ser Ala Leu Pro Glu Asp Ile His Ser
 1345 1350 1355 1360
 Thr Phe Asn Ser Leu Thr Leu Gln Phe Asp Ser Asp Phe Phe Ile Ser
 1365 1370 1375
 Lys Ser Gly Phe Ser Ile Gln Phe Ser Thr Ser Ile Ala Ala Thr Cys
 1380 1385 1390
 Asn Asp Pro Gly Met Pro Gln Asn Gly Thr Arg Tyr Gly Asp Ser Arg
 1395 1400 1405
 Glu Ala Gly Asp Thr Val Thr Phe Gln Cys Asp Pro Gly Tyr Gln Leu
 1410 1415 1420
 Gln Gly Gln Ala Lys Ile Thr Cys Val Gln Leu Asn Asn Arg Phe Phe
 1425 1430 1435 1440
 Trp Gln Pro Asp Pro Pro Thr Cys Ile Ala Ala Cys Gly Gly Asn Leu
 1445 1450 1455

Thr Gly Pro Ala Gly Val Ile Leu Ser Pro Asn Tyr Pro Gln Pro Tyr
 1460 1465 1470

 Pro Pro Gly Lys Glu Cys Asp Trp Arg Val Lys Val Asn Pro Asp Phe
 1475 1480 1485

 Val Ile Ala Leu Ile Phe Lys Ser Phe Asn Met Glu Pro Ser Tyr Asp
 1490 1495 1500

 Phe Leu His Ile Tyr Glu Gly Glu Asp Ser Asn Ser Pro Leu Ile Gly
 1505 1510 1515 1520

 Ser Tyr Gln Gly Ser Gln Ala Pro Glu Arg Ile Glu Ser Ser Gly Asn
 1525 1530 1535

 Ser Leu Phe Leu Ala Phe Arg Ser Asp Ala Ser Val Gly Leu Ser Gly
 1540 1545 1550

 Phe Ala Ile Glu Phe Lys Glu Lys Pro Arg Glu Ala Cys Phe Asp Pro
 1555 1560 1565

 Gly Asn Ile Met Asn Gly Thr Arg Val Gly Thr Asp Phe Lys Leu Gly
 1570 1575 1580

 Ser Thr Ile Thr Tyr Gln Cys Asp Ser Gly Tyr Lys Ile Leu Asp Pro
 1585 1590 1595 1600

 Ser Ser Ile Thr Cys Val Ile Gly Ala Asp Gly Lys Pro Ser Trp Asp
 1605 1610 1615

 Gln Val Leu Pro Ser Cys Asn Ala Pro Cys Gly Gly Gln Tyr Thr Gly
 1620 1625 1630

 Ser Glu Gly Val Val Leu Ser Pro Asn Tyr Pro His Asn Tyr Thr Ala
 1635 1640 1645

 Gly Gln Ile Cys Leu Tyr Ser Ile Thr Val Pro Lys Glu Phe Val Val
 1650 1655 1660

 Phe Gly Gln Phe Ala Tyr Phe Gln Thr Ala Leu Asn Asp Leu Ala Glu
 1665 1670 1675 1680

 Leu Phe Asp Gly Thr His Ala Gln Ala Arg Leu Leu Ser Ser Leu Ser
 1685 1690 1695

 Gly Ser His Ser Gly Glu Thr Leu Pro Leu Ala Thr Ser Asn Gln Ile
 1700 1705 1710

 Leu Leu Arg Phe Ser Ala Lys Ser Gly Ala Ser Ala Arg Gly Phe His
 1715 1720 1725

 Phe Val Tyr Gln Ala Val Pro Arg Thr Ser Asp Thr Gln Cys Ser Ser
 1730 1735 1740

 Val Pro Glu Pro Arg Tyr Gly Arg Arg Ile Gly Ser Glu Phe Ser Ala
 1745 1750 1755 1760

Gly Ser Ile Val Arg Phe Glu Cys Asn Pro Gly Tyr Leu Leu Gln Gly
 1765 1770 1775
 Ser Thr Ala Leu His Cys Gln Ser Val Pro Asn Ala Leu Ala Gln Trp
 1780 1785 1790
 Asn Asp Thr Ile Pro Ser Cys Val Val Pro Cys Ser Gly Asn Phe Thr
 1795 1800 1805
 Gln Arg Arg Gly Thr Ile Leu Ser Pro Gly Tyr Pro Glu Pro Tyr Gly
 1810 1815 1820
 Asn Asn Leu Asn Cys Ile Trp Lys Ile Ile Val Thr Glu Gly Ser Gly
 1825 1830 1835 1840
 Ile Gln Asp Pro Ser Asp Gln Phe Cys His Gly Ala Glu Leu Gly Leu
 1845 1850 1855
 Pro Phe Glu Ile His Asp Gly Gly Asp Val Thr Ala Pro Arg Leu Gly
 1860 1865 1870
 Ser Phe Ser Gly Thr Thr Val Pro Ala Leu Leu Asn Ser Thr Ser Asn
 1875 1880 1885
 Gln Leu Tyr Leu His Phe Gln Ser Asp Ile Ser Val Ala Ala Ala Gly
 1890 1895 1900
 Phe His Leu Glu Tyr Lys Thr Val Gly Leu Ala Ala Cys Gln Glu Pro
 1905 1910 1915 1920
 Ala Leu Pro Ser Asn Ser Ile Lys Ile Gly Asp Arg Tyr Met Val Asn
 1925 1930 1935
 Asp Val Leu Ser Phe Gln Cys Glu Pro Gly Tyr Thr Leu Gln Gly Arg
 1940 1945 1950
 Ser His Ile Ser Cys Met Pro Gly Thr Val Arg Arg Trp Asn Tyr Pro
 1955 1960 1965
 Ser Pro Leu Cys Ile Ala Thr Cys Gly Gly Thr Leu Ser Thr Leu Gly
 1970 1975 1980
 Gly Val Ile Leu Ser Pro Gly Phe Pro Gly Ser Tyr Pro Asn Asn Leu
 1985 1990 1995 2000
 Asp Cys Thr Trp Arg Ile Ser Leu Pro Ile Gly Tyr Gly Ala His Ile
 2005 2010 2015
 Gln Phe Leu Asn Phe Ser Thr Glu Ala Asn His Asp Phe Leu Glu Ile
 2020 2025 2030
 Gln Asn Gly Pro Tyr His Thr Ser Pro Met Ile Gly Gln Phe Ser Gly
 2035 2040 2045
 Thr Asp Leu Pro Ala Ala Leu Leu Ser Thr Thr His Glu Thr Leu Ile
 2050 2055 2060

His Phe Tyr Ser Asp His Ser Gln Asn Arg Gln Gly Phe Lys Leu Ala
 2065 2070 2075 2080
 Tyr Gln Ala Tyr Glu Leu Gln Asn Cys Pro Asp Pro Pro Phe Gln
 2085 2090 2095
 Asn Gly Tyr Met Ile Asn Ser Asp Tyr Ser Val Gly Gln Ser Val Ser
 2100 2105 2110
 Phe Glu Cys Tyr Pro Gly Tyr Ile Leu Ile Gly His Pro Val Leu Thr
 2115 2120 2125
 Cys Gln His Gly Ile Asn Arg Asn Trp Asn Tyr Pro Phe Pro Arg Cys
 2130 2135 2140
 Asp Ala Pro Cys Gly Tyr Asn Val Thr Ser Gln Asn Gly Thr Ile Tyr
 2145 2150 2155 2160
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 Pro His Gly Tyr Pro Asn Tyr Ala Asn Cys Thr Trp Ile Ile Ile Thr
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 Gly Glu Arg Asn Arg Ile Gln Leu Ser Phe His Thr Phe Ala Leu Glu
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 Glu Asn Phe Asp Ile Leu Ser Val Tyr Asp Gly Gln Pro Gln Gln Gly
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 Asn Leu Lys Val Arg Leu Ser Gly Phe Gln Leu Pro Ser Ser Ile Val
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 Ser Thr Gly Ser Ile Leu Thr Leu Trp Phe Thr Thr Asp Phe Ala Val
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 Ser Ala Gln Gly Phe Lys Ala Leu Tyr Glu Val Leu Pro Ser His Thr
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 Glu Gly Tyr Asp Phe Leu Glu Ile Ser Gly Thr Glu Ala Pro Ser Ile
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 Gly Ala Asn Val Gln Phe Ser Cys Glu Asp Asn Tyr Val Leu Gln Gly
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 Ser Lys Ser Ile Thr Cys Gln Arg Val Thr Glu Thr Leu Ala Ala Trp
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 Arg Gly Pro Ser Gly Val Ile Thr Ser Pro Asn Tyr Pro Val Gln Tyr
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 Glu Asp Asn Ala His Cys Val Trp Val Ile Thr Thr Thr Asp Pro Asp
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 Lys Val Ile Lys Leu Ala Phe Glu Glu Phe Glu Leu Glu Arg Gly Tyr
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 Asp Thr Leu Thr Val Gly Asp Ala Gly Lys Val Gly Asp Thr Arg Ser
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 Val Leu Tyr Val Leu Thr Gly Ser Ser Val Pro Asp Leu Ile Val Ser
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 Met Ser Asn Gln Met Trp Leu His Leu Gln Ser Asp Asp Ser Ile Gly
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 Ser Pro Gly Phe Lys Ala Val Tyr Gln Glu Ile Glu Lys Gly Gly Cys
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 Met Asn Cys Val Trp Leu Ile Ile Ser Glu Pro Gly Ser Arg Ile His
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 Leu Ile Phe Asn Asp Phe Asp Val Glu Pro Gln Phe Asp Phe Leu Ala
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 660 665 670
 Leu Glu Phe Gln Ser Asp His Ser Thr Thr Gly Arg Gly Phe Asn Ile
 675 680 685
 Thr Tyr Thr Thr Phe Gly Gln Asn Glu Cys His Asp Pro Gly Ile Pro
 690 695 700
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 Ser Phe His Cys Asp Asp Gly Phe Val Lys Thr Gln Gly Ser Glu Ser
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 Pro Arg Cys Glu Ala Pro Cys Gly Gly His Leu Thr Ala Ser Ser Gly
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 Val Ile Leu Pro Pro Gly Trp Pro Gly Tyr Tyr Lys Asp Ser Leu His
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 Val Cys Glu Arg Asn His Gln Trp Asn His Ala Leu Pro Ser Cys Asp
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 Ala Leu Cys Gly Gly Tyr Ile Gln Gly Lys Ser Gly Thr Val Leu Ser
 930 935 940

Pro Gly Phe Pro Asp Phe Tyr Pro Asn Ser Leu Asn Cys Thr Trp Thr
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 2930 2935 2940
 Ser Tyr Ser Cys Met Asp Gly Tyr Gln Leu Ser His Ser Ala Ile Leu
 2945 2950 2955 2960
 Ser Cys Glu Gly Arg Gly Val Trp Lys Gly Glu Ile Pro Gln Cys Leu
 2965 2970 2975
 Pro Val Phe Cys Gly Asp Pro Gly Ile Pro Ala Glu Gly Arg Leu Ser
 2980 2985 2990
 Gly Lys Ser Phe Thr Tyr Lys Ser Glu Val Phe Phe Gln Cys Lys Ser
 2995 3000 3005
 Pro Phe Ile Leu Val Gly Ser Ser Arg Arg Val Cys Gln Ala Asp Gly
 3010 3015 3020
 Thr Trp Ser Gly Ile Gln Pro Thr Cys Ile Asp Pro Ala His Asn Thr
 3025 3030 3035 3040
 Cys Pro Asp Pro Gly Thr Pro His Phe Gly Ile Gln Asn Ser Ser Arg
 3045 3050 3055
 Gly Tyr Glu Val Gly Ser Thr Val Phe Phe Arg Cys Arg Lys Gly Tyr
 3060 3065 3070

His Ile Gln Gly Ser Thr Thr Arg Thr Cys Leu Ala Asn Leu Thr Trp
 3075 3080 3085
 Ser Gly Ile Gln Thr Glu Cys Ile Pro His Ala Cys Arg Gln Pro Glu
 3090 3095 3100
 Thr Pro Ala His Ala Asp Val Arg Ala Ile Asp Leu Pro Thr Phe Gly
 3105 3110 3115 3120
 Tyr Thr Leu Val Tyr Thr Cys His Pro Gly Phe Phe Leu Ala Gly Gly
 3125 3130 3135
 Ser Glu His Arg Thr Cys Lys Ala Asp Met Lys Trp Thr Gly Lys Ser
 3140 3145 3150
 Pro Val Cys Lys Ser Lys Gly Val Arg Glu Val Asn Glu Thr Val Thr
 3155 3160 3165
 Lys Thr Pro Val Pro Ser Asp Val Phe Phe Val Asn Ser Leu Trp Lys
 3170 3175 3180
 Gly Tyr Tyr Glu Tyr Leu Gly Lys Arg Gln Pro Ala Thr Leu Thr Val
 3185 3190 3195 3200
 Asp Trp Phe Asn Ala Thr Ser Ser Lys Val Asn Ala Thr Phe Ser Glu
 3205 3210 3215
 Ala Ser Pro Val Glu Leu Lys Leu Thr Gly Ile Tyr Lys Lys Glu Glu
 3220 3225 3230
 Ala His Leu Leu Leu Lys Ala Phe Gln Ile Lys Gly Gln Ala Asp Ile
 3235 3240 3245
 Phe Val Ser Lys Phe Glu Asn Asp Asn Trp Gly Leu Asp Gly Tyr Val
 3250 3255 3260
 Ser Ser Gly Leu Glu Arg Gly Phe Thr Phe Gln Gly Asp Ile His
 3265 3270 3275 3280
 Gly Lys Asp Phe Gly Lys Phe Lys Leu Glu Arg Gln Asp Pro Leu Asn
 3285 3290 3295
 Pro Asp Gln Asp Ser Ser His Tyr His Gly Thr Ser Ser Gly Ser
 3300 3305 3310
 Val Ala Ala Ala Ile Leu Val Pro Phe Phe Ala Leu Ile Leu Ser Gly
 3315 3320 3325
 Phe Ala Phe Tyr Leu Tyr Lys His Arg Thr Arg Pro Lys Val Gln Tyr
 3330 3335 3340
 Asn Gly Tyr Ala Gly His Glu Asn Ser Asn Gly Gln Ala Ser Phe Glu
 3345 3350 3355 3360
 Asn Pro Met Tyr Asp Thr Asn Leu Lys Pro Thr Glu Ala Lys Ala Val
 3365 3370 3375

Arg Phe Asp Thr Thr Leu Asn Thr Val Cys Thr Val Val
3380 3385

<210> 48
<211> 1043
<212> PRT
<213> Homo sapiens

<400> 48
Met Ile Phe His Thr Phe His Leu Glu Ser Ser His Asp Tyr Leu Leu
1 5 10 15
Ile Thr Glu Asp Gly Ser Phe Ser Glu Pro Val Ala Arg Leu Thr Gly
20 25 30
Ser Val Leu Pro His Thr Ile Lys Ala Gly Leu Phe Gly Asn Phe Thr
35 40 45
Ala Gln Leu Arg Phe Ile Ser Asp Phe Ser Ile Ser Tyr Glu Gly Phe
50 55 60
Asn Ile Thr Phe Ser Glu Tyr Asp Leu Glu Pro Cys Asp Asp Pro Gly
65 70 75 80
Val Pro Ala Phe Ser Arg Arg Ile Gly Phe His Phe Gly Val Gly Asp
85 90 95
Ser Leu Thr Phe Ser Cys Phe Leu Gly Tyr Arg Leu Glu Gly Ala Thr
100 105 110
Lys Leu Thr Cys Leu Gly Gly Arg Arg Val Trp Ser Ala Pro Leu
115 120 125
Pro Arg Cys Val Ala Glu Cys Gly Ala Ser Val Lys Gly Asn Glu Gly
130 135 140
Thr Leu Leu Ser Pro Asn Phe Pro Ser Asn Tyr Asp Asn Asn His Glu
145 150 155 160
Cys Ile Tyr Lys Ile Glu Thr Glu Ala Gly Lys Gly Ile His Leu Arg
165 170 175
Thr Arg Ser Phe Gln Leu Phe Glu Gly Asp Thr Leu Lys Val Tyr Asp
180 185 190
Gly Lys Asp Ser Ser Arg Pro Leu Gly Thr Phe Thr Lys Asn Glu
195 200 205
Leu Leu Gly Leu Ile Leu Asn Ser Thr Ser Asn His Leu Trp Leu Glu
210 215 220
Phe Asn Thr Asn Gly Ser Asp Thr Asp Gln Gly Phe Gln Leu Thr Tyr
225 230 235 240
Thr Ser Phe Asp Leu Val Lys Cys Glu Asp Pro Gly Ile Pro Asn Tyr

245	250	255
Gly Tyr Arg Ile Arg Asp Glu Gly His Phe Thr Asp Thr Val Val Leu		
260	265	270
Tyr Ser Cys Asn Pro Gly Tyr Ala Met His Gly Ser Asn Thr Leu Thr		
275	280	285
Cys Leu Ser Gly Asp Arg Arg Val Trp Asp Lys Pro Leu Pro Ser Cys		
290	295	300
Ile Ala Glu Cys Gly Gly Gln Ile His Ala Ala Thr Ser Gly Arg Ile		
305	310	315
Leu Ser Pro Gly Tyr Pro Ala Pro Tyr Asp Asn Asn Leu His Cys Thr		
325	330	335
Trp Ile Ile Glu Ala Asp Pro Gly Lys Thr Ile Ser Leu His Phe Ile		
340	345	350
Val Phe Asp Thr Glu Met Ala His Asp Ile Leu Lys Val Trp Asp Gly		
355	360	365
Pro Val Asp Ser Asp Ile Leu Leu Lys Glu Trp Ser Gly Ser Ala Leu		
370	375	380
Pro Glu Asp Ile His Ser Thr Phe Asn Ser Leu Thr Leu Gln Phe Asp		
385	390	400
Ser Asp Phe Phe Ile Ser Lys Ser Gly Phe Ser Ile Gln Phe Ser Thr		
405	410	415
Ser Ile Ala Ala Thr Cys Asn Asp Pro Gly Met Pro Gln Asn Gly Thr		
420	425	430
Arg Tyr Gly Asp Ser Arg Glu Ala Gly Asp Thr Val Thr Phe Gln Cys		
435	440	445
Asp Pro Gly Tyr Gln Leu Gln Gly Gln Ala Lys Ile Thr Cys Val Gln		
450	455	460
Leu Asn Asn Arg Phe Phe Trp Gln Pro Asp Pro Pro Thr Cys Ile Ala		
465	470	475
Ala Cys Gly Gly Asn Leu Thr Gly Pro Ala Gly Val Ile Leu Ser Pro		
485	490	495
Asn Tyr Pro Gln Pro Tyr Pro Pro Gly Lys Glu Cys Asp Trp Arg Val		
500	505	510
Lys Val Asn Pro Asp Phe Val Ile Ala Leu Ile Phe Lys Ser Phe Asn		
515	520	525
Met Glu Pro Ser Tyr Asp Phe Leu His Ile Tyr Glu Gly Glu Asp Ser		
530	535	540
Asn Ser Pro Leu Ile Gly Ser Tyr Gln Gly Ser Gln Ala Pro Glu Arg		

545	550	555	560
Ile Glu Ser Ser Gly Asn Ser Leu Phe Leu Ala Phe Arg Ser Asp Ala			
565	570	575	
Ser Val Gly Leu Ser Gly Phe Ala Ile Glu Phe Lys Glu Lys Pro Arg			
580	585	590	
Glu Ala Cys Phe Asp Pro Gly Asn Ile Met Asn Gly Thr Arg Val Gly			
595	600	605	
Thr Asp Phe Lys Leu Gly Ser Thr Ile Thr Tyr Gln Cys Asp Ser Gly			
610	615	620	
Tyr Lys Ile Leu Asp Pro Ser Ser Ile Thr Cys Val Ile Gly Ala Asp			
625	630	635	640
Gly Lys Pro Ser Trp Asp Gln Val Leu Pro Ser Cys Asn Ala Pro Cys			
645	650	655	
Gly Gly Gln Tyr Thr Gly Ser Glu Gly Val Val Leu Ser Pro Asn Tyr			
660	665	670	
Pro His Asn Tyr Thr Ala Gly Gln Ile Cys Leu Tyr Ser Ile Thr Val			
675	680	685	
Pro Lys Glu Phe Val Val Phe Gly Gln Phe Ala Tyr Phe Gln Thr Ala			
690	695	700	
Leu Asn Asp Leu Ala Glu Leu Phe Asp Gly Thr His Ala Gln Ala Arg			
705	710	715	720
Leu Leu Ser Ser Leu Ser Gly Ser His Ser Gly Glu Thr Leu Pro Leu			
725	730	735	
Ala Thr Ser Asn Gln Ile Leu Leu Arg Phe Ser Ala Lys Ser Gly Ala			
740	745	750	
Ser Ala Arg Gly Phe His Phe Val Tyr Gln Ala Val Pro Arg Thr Ser			
755	760	765	
Asp Thr Gln Cys Ser Ser Val Pro Glu Pro Arg Tyr Gly Arg Arg Ile			
770	775	780	
Gly Ser Glu Phe Ser Ala Gly Ser Ile Val Arg Phe Glu Cys Asn Pro			
785	790	795	800
Gly Tyr Leu Leu Gln Gly Ser Thr Ala Leu His Cys Gln Ser Val Pro			
805	810	815	
Asn Ala Leu Ala Gln Trp Asn Asp Thr Ile Pro Ser Cys Val Val Pro			
820	825	830	
Cys Ser Gly Asn Phe Thr Gln Arg Arg Gly Thr Ile Leu Ser Pro Gly			
835	840	845	
Tyr Pro Glu Pro Tyr Gly Asn Asn Leu Asn Cys Ile Trp Lys Ile Ile			

850	855	860
Val Thr Glu Gly Ser Gly Ile Gln Ile Gln Val Ile Ser Phe Ala Thr		
865	870	875
Glu Gln Asn Trp Asp Ser Leu Glu Ile His Asp Gly Gly Asp Val Thr		
885	890	895
Ala Pro Arg Leu Gly Ser Phe Ser Gly Thr Thr Val Pro Ala Leu Leu		
900	905	910
Asn Ser Thr Ser Asn Gln Leu Tyr Leu His Phe Gln Ser Asp Ile Ser		
915	920	925
Val Ala Ala Ala Gly Phe His Leu Glu Tyr Lys Thr Val Gly Leu Ala		
930	935	940
Ala Cys Gln Glu Pro Ala Leu Pro Ser Asn Ser Ile Lys Ile Gly Asp		
945	950	955
Arg Tyr Met Val Asn Asp Val Leu Ser Phe Gln Cys Glu Pro Gly Tyr		
965	970	975
Thr Leu Gln Gly Arg Ser His Ile Ser Cys Met Pro Gly Thr Val Arg		
980	985	990
Arg Trp Asn Tyr Pro Ser Pro Leu Cys Ile Ala Thr Cys Gly Gly Thr		
995	1000	1005
Leu Ser Thr Leu Gly Gly Val Ile Leu Ser Pro Gly Phe Pro Gly Ser		
1010	1015	1020
Tyr Pro Asn Asn Leu Asp Cys Thr Trp Arg Ile Ser Leu Pro Ile Gly		
1025	1030	1035
Tyr Gly Lys		

<210> 49
 <211> 1048
 <212> PRT
 <213> Homo sapiens

<400> 49		
Gly Lys Gly Val Gln Met Ile Phe His Thr Phe His Leu Glu Ser Ser		
1	5	10
		15
His Asp Tyr Leu Leu Ile Thr Glu Asp Gly Ser Phe Ser Glu Pro Val		
20	25	30
Ala Arg Leu Thr Gly Ser Val Leu Pro His Thr Ile Lys Ala Gly Leu		
35	40	45
Phe Gly Asn Phe Thr Ala Gln Leu Arg Phe Ile Ser Asp Phe Ser Ile		
50	55	60

Ser Tyr Glu Gly Phe Asn Ile Thr Phe Ser Glu Tyr Asp Leu Glu Pro
 65 70 75 80
 Cys Asp Asp Pro Gly Val Pro Ala Phe Ser Arg Arg Ile Gly Phe His
 85 90 95
 Phe Gly Val Gly Asp Ser Leu Thr Phe Ser Cys Phe Leu Gly Tyr Arg
 100 105 110
 Leu Glu Gly Ala Thr Lys Leu Thr Cys Leu Gly Gly Arg Arg Val
 115 120 125
 Trp Ser Ala Pro Leu Pro Arg Cys Val Ala Glu Cys Gly Ala Ser Val
 130 135 140
 Lys Gly Asn Glu Gly Thr Leu Leu Ser Pro Asn Phe Pro Ser Asn Tyr
 145 150 155 160
 Asp Asn Asn His Glu Cys Ile Tyr Lys Ile Glu Thr Glu Ala Gly Lys
 165 170 175
 Gly Ile His Leu Arg Thr Arg Ser Phe Gln Leu Phe Glu Gly Asp Thr
 180 185 190
 Leu Lys Val Tyr Asp Gly Lys Asp Ser Ser Ser Arg Pro Leu Gly Thr
 195 200 205
 Phe Thr Lys Asn Glu Leu Leu Gly Leu Ile Leu Asn Ser Thr Ser Asn
 210 215 220
 His Leu Trp Leu Glu Phe Asn Thr Asn Gly Ser Asp Thr Asp Gln Gly
 225 230 235 240
 Phe Gln Leu Thr Tyr Thr Ser Phe Asp Leu Val Lys Cys Glu Asp Pro
 245 250 255
 Gly Ile Pro Asn Tyr Gly Tyr Arg Ile Arg Asp Glu Gly His Phe Thr
 260 265 270
 Asp Thr Val Val Leu Tyr Ser Cys Asn Pro Gly Tyr Ala Met His Gly
 275 280 285
 Ser Asn Thr Leu Thr Cys Leu Ser Gly Asp Arg Arg Val Trp Asp Lys
 290 295 300
 Pro Leu Pro Ser Cys Ile Ala Glu Cys Gly Gln Ile His Ala Ala
 305 310 315 320
 Thr Ser Gly Arg Ile Leu Ser Pro Gly Tyr Pro Ala Pro Tyr Asp Asn
 325 330 335
 Asn Leu His Cys Thr Trp Ile Ile Glu Ala Asp Pro Gly Lys Thr Ile
 340 345 350
 Ser Leu His Phe Ile Val Phe Asp Thr Glu Met Ala His Asp Ile Leu
 355 360 365

Lys Val Trp Asp Gly Pro Val Asp Ser Asp Ile Leu Leu Lys Glu Trp
 370 375 380

 Ser Gly Ser Ala Leu Pro Glu Asp Ile His Ser Thr Phe Asn Ser Leu
 385 390 395 400

 Thr Leu Gln Phe Asp Ser Asp Phe Phe Ile Ser Lys Ser Gly Phe Ser
 405 410 415

 Ile Gln Phe Ser Thr Ser Ile Ala Ala Thr Cys Asn Asp Pro Gly Met
 420 425 430

 Pro Gln Asn Gly Thr Arg Tyr Gly Asp Ser Arg Glu Ala Gly Asp Thr
 435 440 445

 Val Thr Phe Gln Cys Asp Pro Gly Tyr Gln Leu Gln Gly Gln Ala Lys
 450 455 460

 Ile Thr Cys Val Gln Leu Asn Asn Arg Phe Phe Trp Gln Pro Asp Pro
 465 470 475 480

 Pro Thr Cys Ile Ala Ala Cys Gly Gly Asn Leu Thr Gly Pro Ala Gly
 485 490 495

 Val Ile Leu Ser Pro Asn Tyr Pro Gln Pro Tyr Pro Pro Gly Lys Glu
 500 505 510

 Cys Asp Trp Arg Val Lys Val Asn Pro Asp Phe Val Ile Ala Leu Ile
 515 520 525

 Phe Lys Ser Phe Asn Met Glu Pro Ser Tyr Asp Phe Leu His Ile Tyr
 530 535 540

 Glu Gly Glu Asp Ser Asn Ser Pro Leu Ile Gly Ser Tyr Gln Gly Ser
 545 550 555 560

 Gln Ala Pro Glu Arg Ile Glu Ser Ser Gly Asn Ser Leu Phe Leu Ala
 565 570 575

 Phe Arg Ser Asp Ala Ser Val Gly Leu Ser Gly Phe Ala Ile Glu Phe
 580 585 590

 Lys Glu Lys Pro Arg Glu Ala Cys Phe Asp Pro Gly Asn Ile Met Asn
 595 600 605

 Gly Thr Arg Val Gly Thr Asp Phe Lys Leu Gly Ser Thr Ile Thr Tyr
 610 615 620

 Gln Cys Asp Ser Gly Tyr Lys Ile Leu Asp Pro Ser Ser Ile Thr Cys
 625 630 635 640

 Val Ile Gly Ala Asp Gly Lys Pro Ser Trp Asp Gln Val Leu Pro Ser
 645 650 655

 Cys Asn Ala Pro Cys Gly Gly Gln Tyr Thr Gly Ser Glu Gly Val Val
 660 665 670

Leu Ser Pro Asn Tyr Pro His Asn Tyr Thr Ala Gly Gln Ile Cys Leu
 675 680 685
 Tyr Ser Ile Thr Val Pro Lys Glu Phe Val Val Phe Gly Gln Phe Ala
 690 695 700 720
 Tyr Phe Gln Thr Ala Leu Asn Asp Leu Ala Glu Leu Phe Asp Gly Thr
 705 710 715 720
 His Ala Gln Ala Arg Leu Leu Ser Ser Leu Ser Gly Ser His Ser Gly
 725 730 735
 Glu Thr Leu Pro Leu Ala Thr Ser Asn Gln Ile Leu Leu Arg Phe Ser
 740 745 750
 Ala Lys Ser Gly Ala Ser Ala Arg Gly Phe His Phe Val Tyr Gln Ala
 755 760 765
 Val Pro Arg Thr Ser Asp Thr Gln Cys Ser Ser Val Pro Glu Pro Arg
 770 775 780
 Tyr Gly Arg Arg Ile Gly Ser Glu Phe Ser Ala Gly Ser Ile Val Arg
 785 790 795 800
 Phe Glu Cys Asn Pro Gly Tyr Leu Leu Gln Gly Ser Thr Ala Leu His
 805 810 815
 Cys Gln Ser Val Pro Asn Ala Leu Ala Gln Trp Asn Asp Thr Ile Pro
 820 825 830
 Ser Cys Val Val Pro Cys Ser Gly Asn Phe Thr Gln Arg Arg Gly Thr
 835 840 845
 Ile Leu Ser Pro Gly Tyr Pro Glu Pro Tyr Gly Asn Asn Leu Asn Cys
 850 855 860
 Ile Trp Lys Ile Ile Val Thr Glu Gly Ser Gly Ile Gln Ile Gln Val
 865 870 875 880
 Ile Ser Phe Ala Thr Glu Gln Asn Trp Asp Ser Leu Glu Ile His Asp
 885 890 895
 Gly Gly Asp Val Thr Ala Pro Arg Leu Gly Ser Phe Ser Gly Thr Thr
 900 905 910
 Val Pro Ala Leu Leu Asn Ser Thr Ser Asn Gln Leu Tyr Leu His Phe
 915 920 925
 Gln Ser Asp Ile Ser Val Ala Ala Ala Gly Phe His Leu Glu Tyr Lys
 930 935 940
 Thr Val Gly Leu Ala Ala Cys Gln Glu Pro Ala Leu Pro Ser Asn Ser
 945 950 955 960
 Ile Lys Ile Gly Asp Arg Tyr Met Val Asn Asp Val Leu Ser Phe Gln
 965 970 975

Cys Glu Pro Gly Tyr Thr Leu Gln Gly Arg Ser His Ile Ser Cys Met
980 985 990

Pro Gly Thr Val Arg Arg Trp Asn Tyr Pro Ser Pro Leu Cys Ile Ala
995 1000 1005

Thr Cys Gly Gly Thr Leu Ser Thr Leu Gly Gly Val Ile Leu Ser Pro
1010 1015 1020

Gly Phe Pro Gly Ser Tyr Pro Asn Asn Leu Asp Cys Thr Trp Arg Ile
1025 1030 1035 1040

Ser Leu Pro Ile Gly Tyr Gly Lys
1045

<210> 50

<211> 124

<212> PRT

<213> Homo sapiens

<400> 50

His Glu Gly Ile Ser Asn Pro Thr Ile Lys Asp Asn Gly Thr Phe Ser
1 5 10 15

Cys Ala Val Lys Asn Pro Pro Asp Val His His Asn Ile Pro Met Thr
20 25 30

Glu Leu Thr Val Thr Glu Arg Gly Phe Gly Thr Met Leu Ser Ser Val
35 40 45

Ala Leu Leu Ser Ile Leu Val Phe Val Pro Ser Ala Val Val Val Ala
50 55 60

Leu Leu Leu Val Arg Met Gly Arg Lys Ala Ala Gly Leu Lys Lys Arg
65 70 75 80

Ser Arg Ser Gly Tyr Lys Ser Ser Ile Glu Val Ser Asp Asp Thr
85 90 95

Asp Gln Glu Glu Glu Ala Cys Met Ala Arg Leu Cys Val Arg Cys
100 105 110

Ala Glu Cys Leu Asp Ser Asp Tyr Glu Glu Thr Tyr
115 120

<210> 51

<211> 219

<212> PRT

<213> Homo sapiens

<400> 51

Ile Val Val Tyr Thr Asp Lys Glu Val His Gly Ala Val Gly Ser Gln
1 5 10 15

Val Thr Leu Tyr Cys Ser Phe Trp Ser Ser Glu Trp Val Ser Asp Asp

20	25	30
Leu Ser Phe Thr Trp Arg Tyr Gln Pro Glu Gly Gly Arg Asp Ala Ile		
35	40	45
Ser Ile Phe His Tyr Ala Lys Gly Gln Pro Tyr Ile Asp Glu Val Gly		
50	55	60
Thr Phe Lys Glu Arg Ile Gln Trp Val Gly Asp Pro His Arg Lys Asp		
65	70	75
Gly Ser Ile Val Ile His Asn Leu Asp Tyr Gly Asp Asn Gly Thr Phe		
85	90	95
Thr Cys Asp Val Lys Asn Pro Pro Asp Ile Val Gly Lys Thr Ser Gln		
100	105	110
Val Thr Leu Tyr Val Phe Glu Lys Val Pro Thr Arg Tyr Gly Val Val		
115	120	125
Leu Gly Ala Val Ile Gly Gly Val Leu Gly Val Val Leu Leu Ala Leu		
130	135	140
Leu Leu Phe Tyr Leu Ile Arg Tyr Cys Trp Leu Arg Arg Gln Ala Ala		
145	150	155
Leu Gln Arg Arg Leu His Ala Met Glu Lys Gly Lys Leu His Lys Thr		
165	170	175
Ala Lys Asp Ala Ser Lys Arg Gly Arg Gln Thr Pro Val Leu Tyr Ala		
180	185	190
Met Leu Asp His Ser Arg Ser Thr Lys Ala Ala Ser Glu Lys Lys Thr		
195	200	205
Lys Gly Leu Gly Glu Ser Arg Lys Asp Lys Lys		
210	215	

<210> 52
 <211> 251
 <212> PRT
 <213> Homo sapiens

<400> 52		
Met Ala Pro Gly Ala Pro Ser Ser Ser Pro Ser Pro Ile Leu Ala Val		
1	5	10
20	25	30
Leu Leu Phe Ser Ser Leu Val Leu Ser Pro Ala Gln Ala Ile Val Val		
Tyr Thr Asp Arg Glu Val His Gly Ala Val Gly Ser Arg Val Thr Leu		
35	40	45
His Cys Ser Phe Trp Ser Ser Glu Trp Val Ser Asp Asp Ile Ser Phe		
50	55	60

Thr Trp Arg Tyr Gln Pro Glu Gly Gly Arg Asp Ala Ile Ser Ile Phe
 65 70 75 80
 His Tyr Ala Lys Gly Gln Pro Tyr Ile Asp Glu Val Gly Thr Phe Lys
 85 90 95
 Glu Arg Ile Gln Trp Val Gly Asp Pro Arg Trp Lys Asp Gly Ser Ile
 100 105 110
 Val Ile His Asn Leu Asp Tyr Ser Asp Asn Gly Thr Phe Thr Cys Asp
 115 120 125
 Val Lys Asn Pro Pro Asp Ile Val Gly Lys Thr Ser Gln Val Thr Leu
 130 135 140
 Tyr Val Phe Glu Lys Val Pro Thr Arg Tyr Gly Val Val Leu Gly Ala
 145 150 155 160
 Val Ile Gly Gly Val Leu Gly Val Val Leu Leu Leu Leu Leu Phe
 165 170 175
 Tyr Val Val Arg Tyr Cys Trp Leu Arg Arg Gln Ala Ala Leu Gln Arg
 180 185 190
 Arg Leu Ser Ala Met Glu Lys Gly Lys Leu His Lys Pro Gly Lys Asp
 195 200 205
 Ala Ser Lys Arg Gly Arg Gln Thr Pro Val Leu Tyr Ala Gln Cys Trp
 210 215 220
 Thr Thr Ala Glu Ala Pro Lys Leu Ser Val Arg Arg Arg Pro Arg Gly
 225 230 235 240
 Trp Gly Ser Leu Ala Arg Ile Arg Asn Ser Gly
 ,245 250

<210> 53
 <211> 258
 <212> PRT
 <213> Homo sapiens

<400> 53
 Met Leu Arg Ala Pro Ala Pro Ala Pro Ala Met Ala Pro Gly Ala Pro
 1 5 10 15
 Ser Ser Ser Pro Ser Pro Ile Leu Ala Val Leu Leu Phe Ser Ser Leu
 20 25 30
 Val Leu Ser Pro Ala Gln Ala Ile Val Val Tyr Thr Asp Arg Glu Val
 35 40 45
 His Gly Ala Val Gly Ser Arg Val Thr Leu His Cys Ser Phe Trp Ser
 50 55 60
 Ser Glu Trp Val Ser Asp Asp Ile Ser Phe Thr Trp Arg Tyr Gln Pro
 65 70 75 80

Glu	Gly	Gly	Arg	Asp	Ala	Ile	Ser	Ile	Phe	His	Tyr	Ala	Lys	Gly	Gln
					85				90				95		
Pro	Tyr	Ile	Asp	Glu	Val	Gly	Thr	Phe	Lys	Glu	Arg	Ile	Gln	Trp	Val
					100			105					110		
Gly	Asp	Pro	Arg	Trp	Lys	Asp	Gly	Ser	Ile	Val	Ile	His	Asn	Leu	Asp
					115			120				125			
Tyr	Ser	Asp	Asn	Gly	Thr	Phe	Thr	Cys	Asp	Val	Lys	Asn	Pro	Pro	Asp
					130		135			140					
Ile	Val	Gly	Lys	Thr	Ser	Gln	Val	Thr	Leu	Tyr	Val	Phe	Glu	Lys	Val
					145		150		155			160			
Pro	Thr	Arg	Tyr	Gly	Val	Val	Leu	Gly	Ala	Val	Ile	Gly	Gly	Val	Leu
					165			170			175				
Gly	Val	Val	Leu	Leu	Leu	Leu	Leu	Leu	Phe	Tyr	Val	Val	Arg	Tyr	Cys
					180			185			190				
Trp	Leu	Arg	Arg	Gln	Ala	Ala	Leu	Gln	Arg	Arg	Leu	Ser	Ala	Met	Glu
					195		200			205					
Lys	Gly	Lys	Leu	His	Lys	Pro	Gly	Lys	Asp	Ala	Ser	Lys	Arg	Gly	Arg
					210		215		220						
Gln	Thr	Pro	Val	Leu	Tyr	Ala	Met	Leu	Asp	His	Ser	Arg	Ser	Thr	Lys
					225		230		235			240			
Ala	Val	Ser	Glu	Lys	Lys	Ala	Lys	Gly	Leu	Gly	Glu	Ser	Arg	Lys	Asp
					245			250			255				

Lys Lys

<210>	54														
<211>	248														
<212>	PRT														
<213>	Homo sapiens														
<400>	54														
Met	Ala	Pro	Gly	Ala	Pro	Ser	Ser	Ser	Pro	Ser	Pro	Ile	Leu	Ala	Val
1					5				10				15		
Leu	Leu	Phe	Ser	Ser	Leu	Val	Leu	Ser	Pro	Ala	Gln	Ala	Ile	Val	Val
					20			25				30			
Tyr	Thr	Asp	Arg	Glu	Val	His	Gly	Ala	Val	Gly	Ser	Arg	Val	Thr	Leu
					35			40			45				
His	Cys	Ser	Phe	Trp	Ser	Ser	Glu	Trp	Val	Ser	Asp	Asp	Ile	Ser	Phe
					50		55			60					
Thr	Trp	Arg	Tyr	Gln	Pro	Glu	Gly	Gly	Arg	Asp	Ala	Ile	Ser	Ile	Phe

65	70	75	80
His Tyr Ala Lys Gly Gln Pro Tyr Ile Asp Glu Val Gly Thr Phe Lys			
85		90	95
Glu Arg Ile Gln Trp Val Gly Asp Pro Arg Trp Lys Asp Gly Ser Ile			
100		105	110
Val Ile His Asn Leu Asp Tyr Ser Asp Asn Gly Thr Phe Thr Cys Asp			
115		120	125
Val Lys Asn Pro Pro Asp Ile Val Gly Lys Thr Ser Gln Val Thr Leu			
130		135	140
Tyr Val Phe Glu Lys Val Pro Thr Arg Tyr Gly Val Val Leu Gly Ala			
145		150	160
Val Ile Gly Gly Val Leu Gly Val Val Leu Leu Leu Leu Leu Phe			
165		170	175
Tyr Val Val Arg Tyr Cys Trp Leu Arg Arg Gln Ala Ala Leu Gln Arg			
180		185	190
Arg Leu Ser Ala Met Glu Lys Gly Lys Leu His Lys Pro Gly Lys Asp			
195		200	205
Ala Ser Lys Arg Gly Arg Gln Thr Pro Val Leu Tyr Ala Met Leu Asp			
210		215	220
His Ser Arg Ser Thr Lys Ala Val Ser Glu Lys Lys Ala Lys Gly Leu			
225		230	240
Gly Glu Ser Arg Lys Asp Lys Lys			
245			
<210> 55			
<211> 2327			
<212> PRT			
<213> Xenopus laevis			
<400> 55			
Met Asn Thr Leu Leu Trp Thr Ile Leu Leu Pro Leu Leu Phe Ser Phe			
1	5	10	15
Cys Val Cys Gln Gln Pro Glu His Gln Asp Leu Glu Met Ser Val Gln			
20		25	30
Tyr Tyr Asp Asp Asn Val Ile Asp Leu Leu Glu Ala Leu Asn Val Thr			
35		40	45
Arg Ser Val Lys Gly Val Thr Lys Ala Lys Gly Ser Asp Pro Ala Ser			
50		55	60
Pro Ala Trp Lys Phe Arg Gln Arg Val Pro His Leu Thr Leu Pro Arg			
65		70	75

Asp Tyr Ser Val Tyr Leu Leu Ser Thr Thr Gln Glu Ser Leu Gly Leu
 85 90 95

 His Phe Val Ala Lys Gln Ala Lys Asn Asn Arg Gly Thr Leu Val Ala
 100 105 110

 Phe Leu Ser Pro Ala Ala Thr Lys Ile Asp Gly Arg Pro Leu Leu Arg
 115 120 125

 Leu Ile Ser Asp Thr His Thr Asp Gln Leu Tyr Phe Glu Tyr Arg Thr
 130 135 140

 Ala Gln Thr Met Glu Pro Ala Ser Leu His Phe Pro Gly Ser Ser Pro
 145 150 155 160

 Phe Ser Gly Ser Gln Trp Ala Arg Val Ala Leu Asn Val Asn Thr His
 165 170 175

 Lys Val Thr Leu Phe Leu Asp Cys Glu Glu Pro Val Val Phe Gly Lys
 180 185 190

 Glu Gly Ala Glu Glu Met Leu Ser Leu Ile Leu Pro Leu Asp Leu Glu
 195 200 205

 Ile Thr Phe Ala Ser Thr Pro Ser Asp Lys Glu Ser Lys Phe Leu Gly
 210 215 220

 Tyr Trp Gln Thr Ala Glu Ile Ser Pro Thr Gly Phe Thr Arg Arg Pro
 225 230 235 240

 Trp His Cys Glu Asn Arg Ser Asp Ser Leu Pro Leu Pro Tyr Ser Leu
 245 250 255

 Ser Gly Glu Arg Gln Met Glu Asp Glu Glu Ile Gln Arg Glu Pro Arg
 260 265 270

 Ala Pro Asp Leu Ser Asp Thr Asp His Tyr Gln Gln Gln Ser Glu
 275 280 285

 Val Pro Ala Gln Leu Leu Ala Lys Asp Asp Arg Leu Gln Arg Leu Glu
 290 295 300

 Glu Ala Val Lys Gly Leu Thr Asn Met Ile Asp Met Ile Lys Ser Gln
 305 310 315 320

 Asn Ala Asp Leu Gln Ala Arg Val Ile Ala Leu Glu Ser Cys Glu Cys
 325 330 335

 Arg Arg Ser Thr Cys Val Trp Glu Asp Lys Glu Tyr Gln Asp Ser Glu
 340 345 350

 Thr Trp Lys Lys Asp Ala Cys Asn Ile Cys Val Cys Val Gly Gly Ser
 355 360 365

 Val Thr Cys Ser Val Arg Lys Asp Trp Pro Gln Cys Leu Gly Cys Phe
 370 375 380

His	Glu	Gly	Arg	Asn	Tyr	Asn	Asn	Lys	Asp	Ile	Phe	Ser	Val	Gly	Pro
385					390					395					400
Cys	Met	Ser	Cys	Ile	Cys	Gln	Ser	Gly	Glu	Val	Ser	Cys	Thr	Pro	Lys
					405				410					415	
Leu	Cys	Pro	Pro	Val	Thr	Cys	Ser	Asp	Pro	Val	Thr	Leu	Pro	Asn	Glut
					420				425					430	
Cys	Cys	Pro	Leu	Cys	Ala	Thr	Gly	Cys	Ser	Asp	Gly	His	Lys	Glu	Gly
					435			440				445			
Asp	Thr	Trp	Arg	Lys	Asp	Thr	Cys	Thr	Thr	Cys	Thr	Cys	Gln	Asn	Gly
					450			455			460				
Thr	Ile	Ser	Cys	Glu	Arg	Glu	Gln	Cys	Pro	Glu	Leu	Thr	Cys	Leu	Lys
					465			470			475				480
Arg	His	Thr	Pro	Pro	Gly	Gln	Cys	Cys	Ala	Lys	Cys	Gln	Gln	Gly	Cys
					485				490					495	
Glu	Tyr	Glu	Gly	Leu	Ile	Tyr	Arg	Asn	Gly	Asp	Tyr	Phe	Leu	Ser	Gln
					500				505					510	
Ser	Asn	Pro	Cys	Val	Asn	Cys	Ser	Cys	Leu	Asn	Asn	Leu	Val	Arg	Cys
					515			520				525			
Leu	Pro	Val	Gln	Cys	Pro	Leu	Pro	Ala	Cys	Thr	Asn	Pro	Val	Pro	Ile
					530			535			540				
Pro	Gly	Gln	Cys	Cys	Pro	Ser	Cys	Pro	Val	Cys	Glu	Leu	Asp	Gly	His
					545			550			555				560
Pro	Leu	Ile	Pro	Gly	Gln	Asn	Val	Thr	Thr	Lys	Asp	Gly	Cys	Arg	Leu
					565				570					575	
Cys	Ser	Cys	Gln	Asp	Gly	Lys	Val	Gln	Cys	Thr	Glu	Ser	Val	Gln	Cys
					580				585					590	
Pro	His	Ile	Cys	Thr	His	Gly	Val	Arg	Ser	Asn	Ser	Cys	Cys	Leu	Asp
					595			600			605				
Cys	Ser	Ala	Cys	Glu	Met	His	Gly	Asp	Ile	Ile	Pro	Asn	Gly	Leu	Thr
					610			615			620				
Phe	Gln	Gly	Asn	Met	Asp	Pro	Cys	Glu	Ser	Cys	Thr	Cys	Gln	Asp	Gly
					625			630			635				640
Asn	Val	His	Cys	Val	Arg	Val	Ser	Cys	Pro	Glu	Leu	Ser	Cys	Val	Leu
					645				650					655	
His	Glu	Lys	Ile	Pro	Gly	Glu	Cys	Cys	Ser	Gln	Cys	Ser	Cys	Met	
					660				665					670	
Asp	Gly	Thr	Val	Lys	Arg	Lys	His	Gly	Glu	Glu	Trp	Lys	Pro	Gln	Gly
					675				680					685	

Asp Pro Cys Gln Ser Cys Arg Cys Leu Glu Gly Arg Val Gln Cys Arg
 690 695 700
 Lys Arg His Cys Ala Ala Leu Cys Arg Asn Pro Leu Pro Pro Arg Pro
 705 710 715 720
 Gly Thr Cys Cys Pro Met Cys Asp Gly Cys Leu Tyr Asn Gly Arg Ser
 725 730 735
 Tyr Leu Asn Gly Gln Pro Val Arg Ser Thr Asp Gln Cys Asn Arg Cys
 740 745 750
 Phe Cys Glu Asn Gly Asn Val Gln Cys Glu Pro Ile Ala Cys Pro Gln
 755 760 765
 Ala Pro Cys Arg Asn Pro Val Arg Arg Thr Gly Glu Cys Cys Pro Arg
 770 775 780
 Cys Glu Gly Cys Glu Tyr Asp Ser Arg His Phe Ala Glu Gly Val Val
 785 790 795 800
 Phe Thr Thr Ala His Asp Pro Cys Leu Gln Cys Thr Cys Leu Ser Gly
 805 810 815
 Glu Val Ser Cys Glu His Leu Asp Arg Lys Cys Pro Pro Ser Gln Cys
 820 825 830
 Ser His Pro Gly Lys Ala Ala Gly Gln Cys Cys Pro Ser Cys Asp Val
 835 840 845
 Cys Asp Phe Glu Gly Ile Leu Tyr Thr Asp Arg Gln Thr Phe Gln Pro
 850 855 860
 Pro Gly His Gly Pro Cys Leu Lys Cys Phe Cys Thr Ile Gly Asn Val
 865 870 875 880
 Arg Cys Val Glu Glu Thr Cys Pro Pro Ala Pro Cys Pro Asn Pro Val
 885 890 895
 Arg Asp Pro Glu Gln Cys Cys Pro Val Cys Lys Val Cys Val Gln Asp
 900 905 910
 Gly Val Glu Phe Leu Glu Gly Ile Glu Trp Glu Leu Asp Gly Asn Pro
 915 920 925
 Cys Ser Ser Cys Thr Cys Arg Asn Gly Asp Thr Val Cys Gly Val Ser
 930 935 940
 Glu Cys Pro Pro Val Ser Cys Leu His Pro Thr Arg Arg Glu Gly Glu
 945 950 955 960
 Cys Cys Pro Val Cys Asp Ser Cys Ser Tyr Asn Gln Arg Leu Tyr Ser
 965 970 975
 Asn Glu Gln Ile Phe Thr Asp Pro Asp Asn Pro Cys Gln Asp Cys Gln
 980 985 990

Cys Lys Asp Gly Thr Val Gln Cys Ser Ser Ile Val Cys Pro Pro Val
 995 1000 1005
 Leu Cys Thr Ile Pro Glu Arg Thr Pro Gly Gln Cys Cys Ala Lys Cys
 1010 1015 1020
 Pro Asp Cys Arg Tyr Gln Asp Gln Ile Phe Leu Glu Gly Glu Gln Phe
 1025 1030 1035 1040
 Ser Asn Pro Leu Asn Gln Cys Gln Glu Cys Trp Cys Arg Asp Gly His
 1045 1050 1055
 Val Thr Cys Thr Asp Arg Gly Cys Thr Gly Ala Leu Cys Ser Tyr Pro
 1060 1065 1070
 Leu Pro Gly Thr Cys Cys Gln Asn Asn Cys Asn Gly Cys Asn Tyr Ala
 1075 1080 1085
 Gly Lys Glu Tyr Pro Asn Gly Ala Asp Phe Pro His Pro Thr Asp Lys
 1090 1095 1100
 Cys Arg Gln Cys His Cys Ile Asn Gly Asn Val Gln Cys Leu Ala Gln
 1105 1110 1115 1120
 Arg Cys Pro Pro Leu Leu Cys Ala Glu Pro Phe Pro Val Pro Gly Glu
 1125 1130 1135
 Cys Cys Pro Gln Cys Pro Val Pro Pro Ala Asp Cys Pro Tyr Ser Gly
 1140 1145 1150
 Val Thr Tyr Arg His Met Gln Arg Phe Tyr Asp Pro Ser Asp Lys Cys
 1155 1160 1165
 Arg Asp Cys Ile Cys Asn Asn Gly Thr Val Thr Cys Gln Arg Lys Pro
 1170 1175 1180
 Cys Ala Pro Thr Pro Cys Leu His Pro Leu Gln Gly Asp Cys Cys Arg
 1185 1190 1195 1200
 Ser Cys Asp Gly Cys Leu Met Ser Gly Lys Glu Leu Ala Asn Gly Glu
 1205 1210 1215
 Gln Phe Pro Gln Pro Ser Asp Pro Cys Ser Val Cys Val Cys Trp Glu
 1220 1225 1230
 Gly Ser Val Thr Cys Gln Pro Lys Thr Cys Pro Val Leu Asn Cys Pro
 1235 1240 1245
 Phe Pro Ala Pro Gly Gln Cys Cys Lys Glu Cys Gln Asp Cys Gln Tyr
 1250 1255 1260
 Phe Gly Glu Val Tyr Leu Asn Gly Gln Glu Phe Ser Ala Pro Glu Asp
 1265 1270 1275 1280
 Ser Cys Ser Arg Cys Val Cys Ala Asp Gly Phe Val Thr Cys Ser Lys
 1285 1290 1295

Lys Pro Cys Tyr Lys Ala Gly Cys Thr His Pro Ser Thr Pro Pro Gly
 1300 1305 1310
 Lys Cys Cys Pro Val Cys Asp Gly Cys Ser Tyr Asn Gly Asp Ala Leu
 1315 1320 1325
 Ile Asn Ser Gln Ser Val Pro Asp Pro Ser Asn Pro Leu Cys Ser Glu
 1330 1335 1340
 Cys Thr Cys Arg Ala Gly Ser Val Gln Cys Val Arg Lys Leu Cys Gly
 1345 1350 1355 1360
 Pro Thr Ser Cys Pro His Pro Val Thr Gly Pro Cys Asp Cys Pro Ile
 1365 1370 1375
 Cys Gln Gly Cys His Phe Gln Gly His Asn Tyr Ile Asp Gly Glu Val
 1380 1385 1390
 Phe Thr Ser Ala Gln Ser Gln Cys Glu Gln Cys Arg Cys Met Arg Gly
 1395 1400 1405
 His Val Thr Cys Gly Pro Arg Pro Cys Asp Gln Val Thr Cys Pro His
 1410 1415 1420
 Pro Ala Glu Asp Pro Cys Met Cys Pro Val Cys Asp Gly Cys Asn Tyr
 1425 1430 1435 1440
 Ser Gly Arg Asp Cys Thr Asn Gly Glu Ser Phe Pro Asp Pro Glu Asp
 1445 1450 1455
 Glu Cys Ser His Cys Thr Cys Arg Asn Gly Glu Val Ala Cys Ile Ser
 1460 1465 1470
 Val Pro Cys Pro Arg Val Ser Cys Met Tyr Pro Ile Thr Pro Arg Gly
 1475 1480 1485
 Glu Cys Cys Pro Arg Cys Thr Gly Ile Cys Lys His Asn Gly Arg Val
 1490 1495 1500
 Tyr Gln Ser Gly Asp Thr Phe His Pro Pro Gly Asp Leu Cys Thr Lys
 1505 1510 1515 1520
 Cys Ser Cys Gln Asn Glu Met Val Asn Cys Gln Arg Val Arg Cys Ser
 1525 1530 1535
 Gln Glu Cys Ser His Pro Val Leu Ser Pro Ala Ser Ser Cys Cys Pro
 1540 1545 1550
 Val Cys Asp Arg Cys Phe Tyr Glu Asn Arg Glu Tyr Ala Asn His Glu
 1555 1560 1565
 Thr Phe Thr Ser Thr Ser Asp Pro Cys Gln Arg Cys Val Cys Leu Asp
 1570 1575 1580
 Gly Ser Val Thr Cys Thr His Val Val Cys Pro Tyr Val Ser Cys Ala
 1585 1590 1595 1600

Asn Pro Ile Thr Lys Pro Gly Gln Cys Cys Arg Glu Cys Pro Val Cys
 1605 1610 1615

 Arg Tyr Gln Gly Lys Glu Phe Ser Glu Gly Ala His Trp Val Pro His
 1620 1625 1630

 Thr Asp Pro Cys Leu Lys Cys Thr Cys Ser Asn Gly His Val Asp Cys
 1635 1640 1645

 Glu Pro Pro Gln Cys Pro Pro Leu Pro Cys Thr Gln Gln Val Thr Asp
 1650 1655 1660

 Pro Gly Thr Cys Cys Pro Arg Cys Arg Gly Cys Val Tyr Asn Gly Arg
 1665 1670 1675 1680

 Glu Tyr Arg Asp Asn Ser Asn Trp Leu Ser Ser Ser Asp His Cys Met
 1685 1690 1695

 Ser Cys Met Cys Val Asp Gly Val Thr Thr Cys Ser Lys Leu Gln Cys
 1700 1705 1710

 Ile Thr Ser Cys Thr Asn Gln Ile Thr Ile Pro Gly Glu Cys Cys Pro
 1715 1720 1725

 Val Cys Ala Asp Cys Ile Ser Asn Ser Lys Val Tyr Leu Pro Gly Asp
 1730 1735 1740

 Ser Tyr Asn Pro Ser Lys Asp Pro Cys Glu Ile Cys Thr Cys Glu Ser
 1745 1750 1755 1760

 Leu Pro Asn Gly Gln Gln Tyr Arg His Cys Thr Lys Lys Gln Cys Pro
 1765 1770 1775

 Ser Leu Leu Asp Cys Pro Arg Ser Tyr Ile Leu Pro Pro Ala Glu Gly
 1780 1785 1790

 Gln Cys Cys Ser Ser Cys Ala Gln Ala Leu Ser Asn Cys Thr Asn Thr
 1795 1800 1805

 Leu Val Gly Asn Glu Ile Gln Ala Thr Asp Asp Pro Cys Tyr Thr Cys
 1810 1815 1820

 His Cys Lys Asp Leu Thr Trp Val Cys Val His Gln Pro Cys Pro Ala
 1825 1830 1835 1840

 Leu Ser Cys Pro Arg Ser Glu Gln Phe Thr His Ser Gly Ser Cys Cys
 1845 1850 1855

 Pro Val Cys Asn Glu Cys Val Val Glu Ile Glu Gly Arg Arg Val Pro
 1860 1865 1870

 Asp Gly Glu Thr Trp Thr Asp Arg Gln Asp Pro Cys Val Thr Cys Thr
 1875 1880 1885

 Cys Thr Leu Gly His Val Glu Cys Gln Ile Glu Glu Cys Gln Pro Val
 1890 1895 1900

Gln Cys Gln Glu Gly Glu Arg Lys Val Lys Arg Pro Gly Thr Cys Cys
 1905 1910 1915 1920
 His Glu Cys Gln Ala Ser Ala Val Ser Cys Trp Tyr Gln Gly Gln Arg
 1925 1930 1935
 Phe Leu Ser Asn Glu His Trp Gln Val Asp Glu Cys Thr Ala Cys Thr
 1940 1945 1950
 Cys Val Ser Gly Glu Val His Cys His Ser Glu Arg Cys Pro Gln Val
 1955 1960 1965
 Ser Cys Thr Ala Glu Glu Thr Pro Ala Leu Ile Pro Gly Met Cys Cys
 1970 1975 1980
 Pro His Cys Ile Pro Arg Pro Ala Thr Cys Ile Ala Phe Gly Asp Pro
 1985 1990 1995 2000
 His Tyr Arg Thr Phe Asp Gly Lys Met Tyr His Phe Gln Gly Ser Cys
 2005 2010 2015
 Thr Tyr Val Leu Ser Glu Asp Cys Glu Gly Asp Phe Ser Ile His
 2020 2025 2030
 Val Thr Asn Asp Asp Arg Gly Leu Arg Gly Val Ser Trp Thr Lys Glu
 2035 2040 2045
 Val Thr Val Leu Ile Gly Asp Ala Val Val Gln Leu Leu Gln Asp Trp
 2050 2055 2060
 Val Val Met Val Asp Tyr Gln Thr Val Glu Leu Pro Phe Leu Lys Glu
 2065 2070 2075 2080
 Pro Tyr Ile Tyr Ile Glu Arg Lys Thr Asn Thr Ile Leu Leu Asn Ser
 2085 2090 2095
 Asn Ile Gly Val Lys Val Gln Trp Asn Gly Arg Ser His Leu Glu Val
 2100 2105 2110
 Ser Val Pro Gly Thr Tyr Arg Asp His Leu Cys Gly Leu Cys Gly Asn
 2115 2120 2125
 Phe Asn Asn Tyr Pro Gln Asp Asp Leu Arg Asp Arg Arg Gly Gln Ile
 2130 2135 2140
 Leu Met Ser Glu Ala Ala Phe Gly Asn Ser Trp Arg Val Gln Ser Ser
 2145 2150 2155 2160
 Asn Asp Ser Ser Ser Cys Trp Asp Gly Gln Asp Val Asp Pro Cys
 2165 2170 2175
 Lys Gln Ala Gly Tyr Arg Ala Arg Lys Glu Ala Asn Gly Arg Cys Lys
 2180 2185 2190
 Leu Leu Lys Ser Ser Val Phe Glu Pro Cys His Arg Val Val Pro Pro
 2195 2200 2205

Glu	Met	Phe	Phe	Ala	Ser	Cys	Val	Tyr	Asp	Leu	Cys	Ala	Cys	Gly	Ala
2210							2215					2220			
Gly	Asp	Glu	Cys	Leu	Cys	Asp	Val	Leu	Glu	Ala	Tyr	Ala	Ser	Glu	Cys
2225							2230				2235				2240
Arg	Glu	Ala	Gly	Val	Ile	Leu	Gln	Trp	Arg	Ser	Pro	Ala	Leu	Cys	Ala
2245								2250				2255			
Val	Gly	Cys	Pro	His	Asp	Arg	Gly	Tyr	Val	Phe	Asp	Glu	Cys	Gly	Pro
2260							2265					2270			
Pro	Cys	Pro	Lys	Thr	Cys	Phe	Asn	Lys	Asp	Val	Pro	Leu	Gly	Val	Leu
2275							2280				2285				
Glu	Ser	His	Cys	Phe	Lys	Pro	Cys	Val	Pro	Gly	Cys	Gln	Cys	Pro	Ala
2290					2295							2300			
Gly	Leu	Val	Glu	His	Glu	Ser	His	Cys	Ile	Pro	Pro	Glu	Ser	Cys	Pro
2305					2310					2315					2320
Lys	Ile	Ile	His	Gly	Asn	Leu									
				2325											

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<211>	751														
<212>	PRT														
<213>	Drosophila melanogaster														
<400>	56														
Met	Cys	Cys	Gln	Ser	Ser	Gly	Gln	Trp	Lys	Phe	Pro	Ala	Gln	Gln	Pro
1									10						15
Arg	Lys	Ser	Leu	Ala	Ser	Arg	Arg	Arg	His	Thr	Gly	Phe	Arg	Pro	Ser
									20	25				30	
Thr	Gln	Leu	Leu	Ile	Leu	Ile	Ala	Val	Leu	Leu	Ala	Leu	Leu	Gln	Gly
									35	40				45	
Arg	Thr	Val	Asp	Ala	Gly	Ala	Gly	Asp	Ser	Leu	Ser	Gly	Val	Arg	Gln
									50	55				60	
Ser	Cys	Ser	Asn	Glu	Gly	Glu	Glu	Val	Gln	Leu	Lys	Asn	Gln	Pro	Gln
									65	70				75	80
Ile	Phe	Thr	Cys	Phe	Lys	Cys	Glu	Cys	Gln	Asn	Gly	Phe	Val	Asn	Cys
									85	90				95	
Arg	Asp	Thr	Cys	Pro	Pro	Val	Asn	Asp	Cys	Tyr	Ile	Leu	Asp	Lys	Ser
									100	105				110	
Asn	Gly	Thr	Cys	Cys	Arg	Arg	Cys	Lys	Gly	Cys	Ser	Phe	Arg	Gly	Met
									115	120				125	
Ser	Tyr	Glu	Ser	Gly	Ser	Glu	Trp	Asn	Asp	Pro	Glu	Asp	Pro	Cys	Lys
									130	135				140	

Thr Tyr Lys Cys Val Ala Thr Val Val Thr Glu Thr Ile Gln Lys Cys
 145 150 155 160
 Tyr Ser Gln Cys Asp Asn Asn Gln Leu Gln Pro Pro Arg Pro Gly Glu
 165 170 175
 Cys Cys Pro Thr Cys Gln Gly Cys Lys Ile Asn Gly Gln Thr Val Ala
 180 185 190
 Glu Gly His Glu Val Asp Ala Ser Ile Asp Asp Arg Cys Leu Val Cys
 195 200 205
 Gln Cys Arg Gly Thr Gln Leu Thr Cys Ser Lys Lys Thr Cys Pro Val
 210 215 220
 Leu Pro Cys Pro Met Ser Lys Gln Ile Lys Arg Pro Asp Glu Cys Cys
 225 230 235 240
 Pro Arg Cys Pro Gln Asn His Ser Phe Leu Pro Val Pro Gly Lys Cys
 245 250 255
 Leu Phe Asn Lys Ser Val Tyr Pro Glu Lys Thr Gln Phe Met Pro Asp
 260 265 270
 Arg Cys Thr Asn Cys Thr Cys Leu Asn Gly Thr Ser Val Cys Gln Arg
 275 280 285
 Pro Thr Cys Pro Ile Leu Glu Cys Ala Pro Glu Phe Gln Glu Pro Asp
 290 295 300
 Gly Cys Cys Pro Arg Cys Ala Val Ala Glu Val Arg Ser Glu Cys Ser
 305 310 315 320
 Leu Asp Gly Ile Val Tyr Gln Asn Asn Glu Thr Trp Asp Met Gly Pro
 325 330 335
 Cys Arg Ser Cys Arg Cys Asn Gly Gly Thr Ile Arg Cys Ala Gln Met
 340 345 350
 Arg Cys Pro Ala Val Lys Cys Arg Ala Asn Glu Glu Leu Lys Gln Pro
 355 360 365
 Pro Gly Glu Cys Cys Gln Arg Cys Val Glu Thr Ala Gly Thr Cys Thr
 370 375 380
 Val Phe Gly Asp Pro His Phe Arg Thr Phe Asp Gly Lys Phe Phe Ser
 385 390 395 400
 Phe Gln Gly Ser Cys Lys Tyr Leu Leu Ala Ser Asp Cys Met Gly Lys
 405 410 415
 Thr Phe His Ile Arg Leu Thr Asn Glu Gly Arg Gly Thr Arg Arg Ser
 420 425 430
 Ser Trp Ala Lys Thr Val Thr Leu Ser Leu Arg Asn Leu Lys Val Asn
 435 440 445

Leu Gly Gln Arg Met Arg Val Lys Val Asn Gly Thr Arg Val Thr Leu
450 455 460

Pro Tyr Phe Val Val Ala Gly Gly Gln Asn Val Thr Ile Glu Arg Leu
465 470 475 480

Ala Asp Gly Gly Ala Val Met Leu Arg Ser Glu Met Gly Leu Thr Leu
485 490 495

Glu Trp Asn Gly Ala Gly Phe Leu Gln Val Ser Val Pro Ala Lys Phe
500 505 510

Lys Lys Arg Leu Cys Gly Leu Cys Gly Asn Phe Asn Gly Ser Ser Arg
515 520 525

Asp Asp Leu Thr Gly Lys Asp Gly Arg Ser His Gly Asp Asp Glu Val
530 535 540

Trp His Phe Ala Asn Ser Trp Lys Val Gly Gly Pro Lys Ser Cys Ser
545 550 555 560

Arg Lys Arg Glu Phe Leu Ala Ala Thr Pro Thr Arg Asp Lys Arg Lys
565 570 575

Ser Asn Phe Tyr Cys His Pro Leu Ser Val Pro Ala Leu Phe Gly Glu
580 585 590

Cys Asn Glu Arg Leu Asn Pro Glu Asn Tyr Lys Ala Ala Cys Arg Met
595 600 605

Asp Val Cys Glu Cys Pro Ser Gly Asp Cys His Cys Asp Ser Phe Ala
610 615 620

Ala Tyr Ala His Glu Cys Arg Arg Leu Gly Val Gln Leu Pro Asp Trp
625 630 635 640

Arg Ser Ala Thr Asn Cys Pro Ala Gly Trp Arg Arg Asn Ala Thr Leu
645 650 655

Ser Ser Phe Lys Gly Asn Gln Phe Tyr Gly Asp Pro Ser Phe Ser Arg
660 665 670

Met Lys Gly Arg Arg Gln Lys Asn His Gln Leu Arg Leu Gln Leu Gln
675 680 685

Gln Glu Gln Gln Gln Arg Ser Lys Gln Gly Gln Lys Gly Arg His Lys
690 695 700

Pro Gly Gly His Asn Gln Leu Asp Arg Gln Gly His Asn Gly Leu Asp
705 710 715 720

Lys Asp Gln Leu Gln Lys Glu Phe Ile Leu Lys His Val Pro Ser Ser
725 730 735

Phe Leu Tyr Pro Arg Ala Pro Asp Arg Thr Pro Pro Pro Leu His
740 745 750

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 <211> 665
 <212> PRT
 <213> Drosophila melanogaster

<400> 57
 Met Arg Ala Lys Asn Gly Phe Val Asn Cys Arg Asp Thr Cys Pro Pro
 1 5 10 15

Val Asn Asp Cys Tyr Ile Leu Asp Lys Ser Asn Gly Thr Cys Cys Arg
 20 25 30

Arg Cys Lys Gly Cys Ser Phe Arg Gly Met Ser Tyr Glu Ser Gly Ser
 35 40 45

Glu Trp Asn Asp Pro Glu Asp Pro Cys Lys Thr Tyr Lys Cys Val Ala
 50 55 60

Thr Val Val Thr Glu Thr Ile Gln Lys Cys Tyr Ser Gln Cys Asp Asn
 65 70 75 80

Asn Gln Leu Gln Pro Pro Arg Pro Gly Glu Cys Cys Pro Thr Cys Gln
 85 90 95

Gly Cys Lys Ile Asn Gly Gln Thr Val Ala Glu Gly His Glu Val Asp
 100 105 110

Ala Ser Ile Asp Asp Arg Cys Leu Val Cys Gln Cys Arg Gly Thr Gln
 115 120 125

Leu Thr Cys Ser Lys Lys Thr Cys Pro Val Leu Pro Cys Pro Met Ser
 130 135 140

Lys Gln Ile Lys Arg Pro Asp Glu Cys Cys Pro Arg Cys Pro Gln Asn
 145 150 155 160

His Ser Phe Leu Pro Val Pro Gly Lys Cys Leu Phe Asn Lys Ser Val
 165 170 175

Tyr Pro Glu Lys Thr Gln Phe Met Pro Asp Arg Cys Thr Asn Cys Thr
 180 185 190

Cys Leu Asn Gly Thr Ser Val Cys Gln Arg Pro Thr Cys Pro Ile Leu
 195 200 205

Glu Cys Ala Pro Glu Phe Gln Glu Pro Asp Gly Cys Cys Pro Arg Cys
 210 215 220

Ala Val Ala Glu Val Arg Ser Glu Cys Ser Leu Asp Gly Ile Val Tyr
 225 230 235 240

Gln Asn Asn Glu Thr Trp Asp Met Gly Pro Cys Arg Ser Cys Arg Cys
 245 250 255

Asn Gly Gly Thr Ile Arg Cys Ala Gln Met Arg Cys Pro Ala Val Lys

	260	265	270
Cys Arg Ala Asn Glu Glu Leu Lys Gln Pro Pro Gly Glu Cys Cys Gln			
275	280	285	
Arg Cys Val Glu Thr Ala Gly Thr Cys Thr Val Phe Gly Asp Pro His			
290	295	300	
Phe Arg Thr Phe Asp Gly Lys Phe Phe Ser Phe Gln Gly Ser Cys Lys			
305	310	315	320
Tyr Leu Leu Ala Ser Asp Cys Met Gly Lys Thr Phe His Ile Arg Leu			
325	330	335	
Thr Asn Glu Gly Arg Gly Thr Arg Arg Ala Ser Trp Ala Lys Thr Val			
340	345	350	
Thr Leu Ser Leu Arg Asn Leu Lys Val Asn Leu Gly Gln Arg Met Arg			
355	360	365	
Val Lys Val Asn Gly Thr Arg Val Thr Leu Pro Tyr Phe Val Val Ala			
370	375	380	
Gly Gly Gln Asn Val Thr Ile Glu Arg Leu Ala Asn Gly Gly Ala Val			
385	390	395	400
Met Leu Arg Ser Glu Met Gly Leu Thr Leu Glu Trp Asn Gly Ala Gly			
405	410	415	
Phe Leu Gln Val Ser Val Pro Ala Lys Phe Lys Lys Arg Leu Cys Gly			
420	425	430	
Leu Cys Gly Asn Phe Asn Gly Ser Ser Arg Asp Asp Leu Thr Gly Lys			
435	440	445	
Asp Gly Arg Ser His Gly Asp Asp Glu Val Trp His Phe Ala Asn Ser			
450	455	460	
Trp Lys Val Gly Gly Pro Lys Ser Cys Ser Arg Lys Arg Glu Phe Leu			
465	470	475	480
Ala Ala Thr Pro Thr Cys Asp Lys Arg Lys Ser Asn Phe Tyr Cys His			
485	490	495	
Pro Leu Ser Val Pro Ala Leu Phe Gly Glu Cys Asn Glu Arg Leu Asn			
500	505	510	
Pro Glu Asn Tyr Lys Ala Ala Cys Arg Met Asp Val Cys Glu Cys Pro			
515	520	525	
Ser Gly Asp Cys His Cys Asp Ser Phe Ala Ala Tyr Ala His Glu Cys			
530	535	540	
Arg Arg Leu Gly Val Gln Leu Pro Asp Trp Arg Ser Ala Thr Asn Cys			
545	550	555	560
Pro Ala Gly Trp Arg Arg Asn Ala Thr Leu Ser Ser Phe Lys Gly Asn			

565	570	575
Gln Phe Tyr Gly Asp Pro Ser Phe Ser Arg Met Lys Gly Arg Arg Gln		
580	585	590
Lys Asn His Gln Leu Arg Leu Gln Leu Gln Glu Gln Gln Gln Arg		
595	600	605
Ser Lys Gln Gly Gln Lys Gly Arg His Lys Pro Gly Gly His Asn Gln		
610	615	620
Leu Asp Arg Gln Gly His Asn Gly Leu Asp Lys Asp Gln Leu Gln Lys		
625	630	635
Glu Phe Ile Leu Lys His Val Pro Ser Ser Phe Leu Tyr Pro Arg Ala		
645	650	655
Pro Asp Arg Thr Pro Pro Pro Leu His		
660	665	
<210> 58		
<211> 462		
<212> PRT		
<213> Mus musculus		
<400> 58		
Met Phe Gly Ser Glu Lys Ser Ile Arg Pro Ser Leu Gly Ser Cys Leu		
1	5	10
		15
Phe Arg Ser Asp Val Tyr Asp Asn Gly Ala Ser Phe Val Tyr Asp Asn		
20	25	30
Cys Thr Val Cys Thr Cys Lys Asp Ser Thr Met Val Cys Lys Lys		
35	40	45
Cys Ser His Pro Gly Val Cys Asn Ser Asp Glu Asp Ala Cys Cys Glu		
50	55	60
Asp Cys Leu Leu Arg Val Pro Pro Glu Asp Ile Lys Val Cys Lys Phe		
65	70	75
		80
Gly Ser Lys Ile Phe Arg Asp Gly Glu Met Trp Ser Ser Val Asn Cys		
85	90	95
Ser Ile Cys Ala Cys Val Lys Gly Lys Thr Glu Cys Arg Lys Lys Gln		
100	105	110
Cys Val Pro Val Ser Ser Cys Pro Gln Gly Lys Ile Leu Asn Arg Lys		
115	120	125
Gly Cys Cys Pro Ile Cys Thr Glu Lys Pro Gly Val Cys Thr Val Phe		
130	135	140
Gly Asp Pro His Tyr Asn Thr Phe Asp Gly Arg Thr Phe Asn Phe Gln		
145	150	155
		160

Gly Thr Cys Gln Tyr Val Leu Thr Lys Asp Cys Ser Ser Pro Ala Ser
 165 170 175

 Pro Phe Gln Val Leu Val Lys Asn Asp Ala Arg Arg Thr Arg Ser Phe
 180 185 190

 Ser Trp Thr Lys Ser Val Glu Leu Met Leu Gly Glu Ser Thr Val Ser
 195 200 205

 Leu Gln Gln His Leu Thr Val Arg Trp Asn Gly Ser Arg Ile Ala Leu
 210 215 220

 Pro Cys His Thr Pro His Phe His Ile Asp Leu Asp Gly Tyr Leu Leu
 225 230 235 240

 Lys Val Thr Thr Arg Ala Gly Leu Glu Ile Ser Trp Asp Gly Asp Ser
 245 250 255

 Phe Val Glu Val Met Ala Ala Pro His Leu Lys Gly Lys Leu Cys Gly
 260 265 270

 Leu Cys Gly Asn Tyr Asn Gly His Lys Arg Asp Asp Leu Ile Gly Gly
 275 280 285

 Asp Gly Asn Phe Lys Phe Asp Val Asp Asp Phe Ala Glu Ser Trp Arg
 290 295 300

 Val Glu Ser Asn Glu Phe Cys Asn Arg Pro Gln Arg Lys Pro Val Pro
 305 310 315 320

 Glu Leu Cys Gln Gly Thr Val Lys Val Lys Leu Arg Ala His Arg Glu
 325 330 335

 Cys Gln Lys Leu Lys Ser Trp Glu Phe Gln Thr Cys His Ser Thr Val
 340 345 350

 Asp Tyr Thr Thr Phe Tyr Arg Ser Cys Val Thr Asp Met Cys Glu Cys
 355 360 365

 Pro Val His Lys Asn Cys Tyr Cys Glu Ser Phe Leu Ala Tyr Thr Arg
 370 375 380

 Ala Cys Gln Arg Glu Gly Ile Lys Val His Trp Glu Pro Gln Gln Ser
 385 390 395 400

 Cys Ala Ala Thr Gln Cys Lys His Gly Ala Val Tyr Asp Thr Cys Gly
 405 410 415

 Pro Gly Cys Val Lys Thr Cys Asp Asn Trp Asn Glu Ile Gly Pro Cys
 420 425 430

 Asn Lys Pro Cys Ile Ala Gly Cys His Cys Pro Ala Asn Leu Val Leu
 435 440 445

 His Lys Gly Arg Cys Ile Lys Pro Val Leu Cys Pro Gln Arg
 450 455 460

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<211> 2601
<212> PRT
<213> Homo sapiens

<400> 59
Met Val Pro Pro Val Trp Thr Leu Leu Leu Val Gly Ala Ala Leu
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Phe Arg Lys Glu Lys Pro Pro Asp Gln Lys Leu Val Val Arg Ser Ser
20 25 30
Arg Asp Asn Tyr Val Leu Thr Gln Cys Asp Phe Glu Asp Asp Ala Lys
35 40 45
Pro Leu Cys Asp Trp Ser Gln Val Ser Ala Asp Asp Glu Asp Trp Val
50 55 60
Arg Ala Ser Gly Pro Ser Pro Thr Gly Ser Thr Gly Ala Pro Gly Gly
65 70 75 80
Tyr Pro Asn Gly Glu Gly Ser Tyr Leu His Met Glu Ser Asn Ser Phe
85 90 95
His Arg Gly Gly Val Ala Arg Leu Leu Ser Pro Asp Leu Trp Glu Gln
100 105 110
Gly Pro Leu Cys Val His Phe Ala His His Met Phe Gly Leu Ser Trp
115 120 125
Gly Ala Gln Leu Arg Leu Leu Leu Ser Gly Glu Glu Gly Arg Arg
130 135 140
Pro Asp Val Leu Trp Lys His Trp Asn Thr Gln Arg Pro Ser Trp Met
145 150 155 160
Leu Thr Thr Val Thr Val Pro Ala Gly Phe Thr Leu Pro Thr Arg Leu
165 170 175
Met Phe Glu Gly Thr Arg Gly Ser Thr Ala Tyr Leu Asp Ile Ala Leu
180 185 190
Asp Ala Leu Ser Ile Arg Arg Gly Ser Cys Asn Arg Val Cys Met Met
195 200 205
Gln Thr Cys Ser Phe Asp Ile Pro Asn Asp Leu Cys Asp Trp Thr Trp
210 215 220
Ile Pro Thr Ala Ser Gly Ala Lys Trp Thr Gln Lys Lys Gly Ser Ser
225 230 235 240
Gly Lys Pro Gly Val Gly Pro Asp Gly Asp Phe Ser Ser Pro Gly Ser
245 250 255
Gly Cys Tyr Met Leu Leu Asp Pro Lys Asn Ala Arg Pro Gly Gln Lys
260 265 270

Ala Val Leu Leu Ser Pro Val Ser Leu Ser Ser Gly Cys Leu Ser Phe
 275 280 285

 Ser Phe His Tyr Ile Leu Arg Gly Gln Ser Pro Gly Ala Ala Leu His
 290 295 300

 Ile Tyr Ala Ser Val Leu Gly Ser Ile Arg Lys His Thr Leu Phe Ser
 305 310 315 320

 Gly Gln Pro Gly Pro Asn Trp Gln Ala Val Ser Val Asn Tyr Thr Ala
 325 330 335

 Val Gly Arg Ile Gln Phe Ala Val Val Gly Val Phe Gly Lys Thr Pro
 340 345 350

 Glu Pro Ala Val Ala Val Asp Ala Thr Ser Ile Ala Pro Cys Gly Glu
 355 360 365

 Gly Phe Pro Gln Cys Asp Phe Glu Asp Asn Ala His Pro Phe Cys Asp
 370 375 380

 Trp Val Gln Thr Ser Gly Asp Gly Gly His Trp Ala Leu Gly His Lys
 385 390 395 400

 Asn Gly Pro Val His Gly Met Gly Pro Ala Gly Gly Phe Pro Asn Ala
 405 410 415

 Gly Gly His Tyr Ile Tyr Leu Glu Ala Asp Glu Phe Ser Gln Ala Gly
 420 425 430

 Gln Ser Val Arg Leu Val Ser Arg Pro Phe Cys Ala Pro Gly Asp Ile
 435 440 445

 Cys Val Glu Phe Ala Tyr His Met Tyr Gly Leu Gly Glu Gly Thr Met
 450 455 460

 Leu Glu Leu Leu Leu Gly Ser Pro Ala Gly Ser Pro Pro Ile Pro Leu
 465 470 475 480

 Trp Lys Arg Val Gly Ser Gln Arg Pro Tyr Trp Gln Asn Thr Ser Val
 485 490 495

 Thr Val Pro Ser Gly His Gln Gln Pro Met Gln Leu Ile Phe Lys Gly
 500 505 510

 Ile Gln Gly Ser Asn Thr Ala Ser Val Val Ala Met Gly Phe Ile Leu
 515 520 525

 Ile Asn Pro Gly Thr Cys Pro Val Lys Val Leu Pro Glu Leu Pro Pro
 530 535 540

 Val Ser Pro Val Ser Ser Thr Gly Pro Ser Glu Thr Thr Gly Leu Thr
 545 550 555 560

 Glu Asn Pro Thr Ile Ser Thr Lys Lys Pro Thr Val Ser Ile Glu Lys
 565 570 575

Pro Ser Val Thr Thr Glu Lys Pro Thr Val Pro Lys Glu Lys Pro Thr
 580 585 590
 Ile Pro Thr Glu Lys Pro Thr Ile Ser Thr Glu Lys Pro Thr Ile Pro
 595 600 605
 Ser Glu Lys Pro Asn Met Pro Ser Glu Lys Pro Thr Ile Pro Ser Glu
 610 615 620
 Lys Pro Thr Ile Leu Thr Glu Lys Pro Thr Ile Pro Ser Glu Lys Pro
 625 630 635 640
 Thr Ile Pro Ser Glu Lys Pro Thr Ile Ser Thr Glu Lys Pro Thr Val
 645 650 655
 Pro Thr Glu Glu Pro Thr Thr Pro Thr Glu Glu Thr Thr Ser Met
 660 665 670
 Glu Glu Pro Val Ile Pro Thr Glu Lys Pro Ser Ile Pro Thr Glu Lys
 675 680 685
 Pro Ser Ile Pro Thr Glu Lys Pro Thr Ile Ser Met Glu Glu Thr Ile
 690 695 700
 Ile Ser Thr Glu Lys Pro Thr Ile Ser Pro Glu Lys Pro Thr Ile Pro
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 Thr Glu Lys Pro Thr Ile Pro Thr Glu Lys Ser Thr Ile Ser Pro Glu
 725 730 735
 Lys Pro Thr Thr Pro Thr Glu Lys Pro Thr Ile Pro Thr Glu Lys Pro
 740 745 750
 Thr Ile Ser Pro Glu Lys Pro Thr Thr Pro Thr Glu Lys Pro Thr Ile
 755 760 765
 Ser Pro Glu Lys Leu Thr Ile Pro Thr Glu Lys Pro Thr Ile Pro Thr
 770 775 780
 Glu Lys Pro Thr Ile Pro Thr Glu Lys Pro Thr Ile Ser Thr Glu Glu
 785 790 795 800
 Pro Thr Thr Pro Thr Glu Glu Thr Thr Ile Ser Thr Glu Lys Pro Ser
 805 810 815
 Ile Pro Met Glu Lys Pro Thr Leu Pro Thr Glu Glu Thr Thr Ser
 820 825 830
 Val Glu Glu Thr Thr Ile Ser Thr Glu Lys Leu Thr Ile Pro Met Glu
 835 840 845
 Lys Pro Thr Ile Ser Thr Glu Lys Pro Thr Ile Pro Thr Glu Lys Pro
 850 855 860
 Thr Ile Ser Pro Glu Lys Leu Thr Ile Pro Thr Glu Lys Leu Thr Ile
 865 870 875 880

Pro Thr Glu Lys Pro Thr Ile Pro Ile Glu Glu Thr Thr Ile Ser Thr
 885 890 895

 Glu Lys Leu Thr Ile Pro Thr Glu Lys Pro Thr Ile Ser Pro Glu Lys
 900 905 910

 Pro Thr Ile Ser Thr Glu Lys Pro Thr Ile Pro Thr Glu Lys Pro Thr
 915 920 925

 Ile Pro Thr Glu Glu Thr Thr Ile Ser Thr Glu Lys Leu Thr Ile Pro
 930 935 940

 Thr Glu Lys Pro Thr Ile Ser Pro Glu Lys Leu Thr Ile Pro Thr Glu
 945 950 955 960

 Lys Pro Thr Ile Ser Thr Glu Lys Pro Thr Ile Pro Thr Glu Lys Leu
 965 970 975

 Thr Ile Pro Thr Glu Lys Pro Thr Ile Pro Thr Glu Lys Pro Thr Ile
 980 985 990

 Pro Thr Glu Lys Leu Thr Ala Leu Arg Pro Pro His Pro Ser Pro Thr
 995 1000 1005

 Ala Thr Gly Leu Ala Ala Leu Val Met Ser Pro His Ala Pro Ser Thr
 1010 1015 1020

 Pro Met Thr Ser Val Ile Leu Gly Thr Thr Thr Ser Arg Ser Ser
 1025 1030 1035 1040

 Thr Glu Arg Cys Pro Pro Asn Ala Arg Tyr Glu Ser Cys Ala Cys Pro
 1045 1050 1055

 Ala Ser Cys Lys Ser Pro Arg Pro Ser Cys Gly Pro Leu Cys Arg Glu
 1060 1065 1070

 Gly Cys Val Cys Asn Pro Gly Phe Leu Phe Ser Asp Asn His Cys Ile
 1075 1080 1085

 Gln Ala Ser Ser Cys Asn Cys Phe Tyr Asn Asn Asp Tyr Tyr Glu Pro
 1090 1095 1100

 Gly Ala Glu Trp Phe Ser Pro Asn Cys Thr Glu His Cys Arg Cys Trp
 1105 1110 1115 1120

 Pro Gly Ser Arg Val Glu Cys Gln Ile Ser Gln Cys Gly Thr His Thr
 1125 1130 1135

 Val Cys Gln Leu Lys Asn Gly Gln Tyr Gly Cys His Pro Tyr Ala Gly
 1140 1145 1150

 Thr Ala Thr Cys Leu Val Tyr Gly Asp Pro His Tyr Val Thr Phe Asp
 1155 1160 1165

 Gly Arg His Phe Gly Phe Met Gly Lys Cys Thr Tyr Ile Leu Ala Gln
 1170 1175 1180

Pro Cys Gly Asn Ser Thr Asp Pro Phe Phe Arg Val Thr Ala Lys Asn
 1185 1190 1195 1200
 Glu Glu Gln Gly Gln Glu Gly Val Ser Cys Leu Ser Lys Val Tyr Val
 1205 1210 1215
 Thr Leu Pro Glu Ser Thr Val Thr Leu Leu Lys Gly Arg Arg Thr Leu
 1220 1225 1230
 Val Gly Gly Gln Gln Val Thr Leu Pro Ala Ile Pro Ser Lys Gly Val
 1235 1240 1245
 Phe Leu Gly Ala Ser Gly Arg Phe Val Glu Leu Gln Thr Glu Phe Gly
 1250 1255 1260
 Leu Arg Val Arg Trp Asp Gly Asp Gln Gln Leu Tyr Val Thr Val Ser
 1265 1270 1275 1280
 Ser Thr Tyr Ser Gly Lys Leu Cys Gly Leu Cys Gly Asn Tyr Asp Gly
 1285 1290 1295
 Asn Ser Asp Asn Asp His Leu Lys Leu Asp Gly Ser Pro Ala Gly Asp
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 Lys Glu Glu Leu Gly Asn Ser Trp Gln Thr Asp Gln Asp Glu Asp Gln
 1315 1320 1325
 Glu Cys Gln Lys Tyr Gln Val Val Asn Ser Pro Ser Cys Asp Ser Ser
 1330 1335 1340
 Leu Gln Ser Ser Met Ser Gly Pro Gly Phe Cys Gly Arg Leu Val Asp
 1345 1350 1355 1360
 Thr His Gly Pro Phe Glu Thr Cys Leu Leu His Val Lys Ala Ala Ser
 1365 1370 1375
 Phe Phe Asp Ser Cys Met Leu Asp Met Cys Gly Phe Gln Gly Leu Gln
 1380 1385 1390
 His Leu Leu Cys Thr His Met Ser Thr Met Thr Thr Cys Gln Asp
 1395 1400 1405
 Ala Gly His Ala Val Lys Pro Trp Arg Glu Pro His Phe Cys Pro Met
 1410 1415 1420
 Ala Cys Pro Pro Asn Ser Lys Tyr Ser Leu Cys Ala Lys Pro Cys Pro
 1425 1430 1435 1440
 Asp Thr Cys His Ser Gly Phe Ser Gly Met Phe Cys Ser Asp Arg Cys
 1445 1450 1455
 Val Glu Ala Cys Glu Cys Asn Pro Gly Phe Val Leu Ser Gly Leu Glu
 1460 1465 1470
 Cys Ile Pro Arg Ser Gln Cys Gly Cys Leu His Pro Ala Gly Ser Tyr
 1475 1480 1485

Phe Lys Val Gly Glu Arg Trp Tyr Lys Pro Gly Cys Lys Glu Leu Cys
 1490 1495 1500
 Val Cys Glu Ser Asn Asn Arg Ile Arg Cys Gln Pro Trp Arg Cys Arg
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 Ala Gln Glu Phe Cys Gly Gln Asp Gly Ile Tyr Gly Cys His Ala
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 Gln Gly Ala Ala Thr Cys Thr Ala Ser Gly Asp Pro His Tyr Leu Thr
 1540 1545 1550
 Phe Asp Gly Ala Leu His His Phe Met Gly Thr Cys Thr Tyr Val Leu
 1555 1560 1565
 Thr Arg Pro Cys Trp Ser Arg Ser Gln Asp Ser Tyr Phe Val Val Ser
 1570 1575 1580
 Ala Thr Asn Glu Asn Arg Gly Gly Ile Leu Glu Val Ser Tyr Ile Lys
 1585 1590 1595 1600
 Ala Val His Val Thr Val Phe Asp Leu Ser Ile Ser Leu Leu Arg Gly
 1605 1610 1615
 Cys Lys Val Met Leu Asn Gly His Arg Val Ala Leu Pro Val Trp Leu
 1620 1625 1630
 Ala Gln Gly Arg Val Thr Ile Arg Leu Ser Ser Asn Leu Val Leu Leu
 1635 1640 1645
 Tyr Thr Asn Phe Gly Leu Gln Val Arg Tyr Asp Gly Ser His Leu Val
 1650 1655 1660
 Glu Val Thr Val Pro Ser Ser Tyr Gly Gly Gln Leu Cys Gly Leu Cys
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 Gly Asn Tyr Asn Asn Ser Leu Asp Asp Asn Leu Arg Pro Asp Arg
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 Lys Leu Ala Gly Asp Ser Met Gln Leu Gly Ala Ala Trp Lys Leu Pro
 1700 1705 1710
 Glu Ser Ser Glu Pro Gly Cys Phe Leu Val Gly Gly Lys Pro Ser Ser
 1715 1720 1725
 Cys Gln Glu Asn Ser Met Ala Asp Ala Trp Asn Lys Asn Cys Ala Ile
 1730 1735 1740
 Leu Ile Asn Pro Gln Gly Pro Phe Ser Gln Cys His Gln Val Val Pro
 1745 1750 1755 1760
 Pro Gln Ser Ser Phe Ala Ser Cys Val His Gly Gln Cys Gly Thr Lys
 1765 1770 1775
 Gly Asp Thr Thr Ala Leu Cys Arg Ser Leu Gln Ala Tyr Ala Ser Leu
 1780 1785 1790

Cys Ala Gln Ala Gly Gln Ala Pro Ala Trp Arg Asn Arg Thr Phe Cys
 1795 1800 1805
 Pro Met Arg Cys Pro Pro Gly Ser Ser Tyr Ser Pro Cys Ser Ser Pro
 1810 1815 1820
 Cys Pro Asp Thr Cys Ser Ser Ile Asn Asn Pro Arg Asp Cys Pro Lys
 1825 1830 1835 1840
 Ala Leu Pro Cys Ala Glu Ser Cys Glu Cys Gln Lys Gly His Ile Leu
 1845 1850 1855
 Ser Gly Thr Ser Cys Val Pro Leu Gly Gln Cys Gly Cys Thr Asp Pro
 1860 1865 1870
 Ala Gly Ser Tyr His Pro Val Gly Glu Arg Trp Tyr Thr Glu Asn Thr
 1875 1880 1885
 Cys Thr Arg Leu Cys Thr Cys Ser Val His Asn Asn Ile Thr Cys Phe
 1890 1895 1900
 Gln Ser Thr Cys Lys Pro Asn Gln Ile Cys Trp Ala Leu Asp Gly Leu
 1905 1910 1915 1920
 Leu Arg Cys Arg Ala Ser Gly Val Gly Val Cys Gln Leu Pro Gly Glu
 1925 1930 1935
 Ser His Tyr Val Ser Phe Asp Gly Ser Asn His Ser Ile Pro Asp Ala
 1940 1945 1950
 Cys Thr Leu Val Leu Val Lys Val Cys His Pro Ala Met Ala Leu Pro
 1955 1960 1965
 Phe Phe Lys Ile Ser Ala Lys His Glu Lys Glu Glu Gly Gly Thr Glu
 1970 1975 1980
 Ala Phe Arg Leu His Glu Val Tyr Ile Asp Ile Tyr Asp Ala Gln Val
 1985 1990 1995 2000
 Thr Leu Gln Lys Gly His Arg Val Leu Ile Asn Ser Lys Gln Val Thr
 2005 2010 2015
 Leu Pro Ala Ile Ser Gln Ile Pro Gly Val Ser Val Lys Ser Ser Ser
 2020 2025 2030
 Ile Tyr Ser Ile Val Asn Ile Lys Ile Gly Val Gln Val Lys Phe Asp
 2035 2040 2045
 Gly Asn His Leu Leu Glu Ile Glu Ile Pro Thr Thr Tyr Tyr Gly Lys
 2050 2055 2060
 Val Cys Gly Met Cys Gly Asn Phe Asn Asp Glu Glu Glu Asp Glu Leu
 2065 2070 2075 2080
 Met Met Pro Ser Asp Glu Val Ala Asn Ser Asp Ser Glu Phe Val Asn
 2085 2090 2095

Ser Trp Lys Asp Lys Asp Ile Asp Pro Ser Cys Gln Ser Leu Leu Val
 2100 2105 2110
 Asp Glu Gln Gln Ile Pro Ala Glu Gln Gln Glu Asn Pro Ser Gly Asn
 2115 2120 2125
 Cys Arg Ala Ala Asp Leu Arg Arg Ala Arg Glu Lys Cys Glu Ala Ala
 2130 2135 2140
 Leu Arg Ala Pro Val Trp Ala Gln Cys Ala Ser Arg Ile Asp Leu Thr
 2145 2150 2155 2160
 Pro Phe Leu Val Asp Cys Ala Asn Thr Leu Cys Glu Phe Gly Gly Leu
 2165 2170 2175
 Tyr Gln Ala Leu Cys Gln Ala Leu Gln Ala Phe Gly Ala Thr Cys Gln
 2180 2185 2190
 Ser Gln Gly Leu Lys Pro Pro Leu Trp Arg Asn Ser Ser Phe Cys Pro
 2195 2200 2205
 Leu Glu Cys Pro Ala Tyr Ser Ser Tyr Thr Asn Cys Leu Pro Ser Cys
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 Ser Pro Ser Cys Trp Asp Leu Asp Gly Arg Cys Glu Gly Ala Lys Val
 2225 2230 2235 2240
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 2245 2250 2255
 Ser Glu Asp Lys Cys Val Pro Arg Ser Gln Cys Gly Cys Lys Asp Ala
 2260 2265 2270
 His Gly Gly Ser Ile Pro Leu Gly Lys Ser Trp Val Ser Ser Gly Cys
 2275 2280 2285
 Thr Glu Lys Cys Val Cys Thr Gly Gly Ala Ile Gln Cys Gly Asp Phe
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 Arg Cys Pro Ser Gly Ser His Cys Gln Leu Thr Ser Asp Asn Ser Asn
 2305 2310 2315 2320
 Ser Asn Cys Val Ser Asp Lys Ser Glu Gln Cys Ser Val Tyr Gly Asp
 2325 2330 2335
 Pro Arg Tyr Leu Thr Phe Asp Gly Phe Ser Tyr Arg Leu Gln Gly Arg
 2340 2345 2350
 Met Thr Tyr Val Leu Ile Lys Thr Val Asp Val Leu Pro Glu Gly Val
 2355 2360 2365
 Glu Pro Leu Leu Val Glu Gly Arg Asn Lys Met Asp Pro Pro Arg Ser
 2370 2375 2380
 Ser Ile Phe Leu Gln Glu Val Ile Thr Thr Val Tyr Gly Tyr Lys Val
 2385 2390 2395 2400

Gln Leu Gln Ala Gly Leu Glu Leu Val Val Asn Asn Gln Lys Met Ala
 2405 2410 2415
 Val Pro Tyr Arg Pro Asn Glu His Leu Arg Val Thr Leu Trp Gly Gln
 2420 2425 2430
 Arg Leu Tyr Leu Val Thr Asp Phe Glu Leu Val Val Ser Phe Gly Gly
 2435 2440 2445
 Arg Lys Asn Ala Val Ile Ser Leu Pro Ser Met Tyr Glu Gly Leu Val
 2450 2455 2460
 Ser Gly Leu Cys Gly Asn Tyr Asp Lys Asn Arg Lys Asn Asp Met Met
 2465 2470 2475 2480
 Leu Pro Ser Gly Ala Leu Thr Gln Asn Leu Asn Thr Phe Gly Asn Ser
 2485 2490 2495
 Trp Glu Val Lys Thr Glu Asp Ala Leu Leu Arg Phe Pro Arg Ala Ile
 2500 2505 2510
 Pro Ala Glu Glu Glu Gly Gln Gly Ala Glu Leu Gly Leu Arg Thr Gly
 2515 2520 2525
 Leu Gln Val Ser Glu Cys Ser Pro Glu Gln Leu Ala Ser Asn Ser Thr
 2530 2535 2540
 Gln Ala Cys Arg Val Leu Ala Asp Pro Gln Gly Pro Phe Ala Ala Cys
 2545 2550 2555 2560
 His Gln Thr Val Ala Pro Glu Pro Phe Gln Glu His Cys Val Leu Asp
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 Leu Cys Ser Ala Gln Asp Pro Arg Glu Gln Glu Glu Leu Arg Cys Gln
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 Val Leu Ser Gly Trp Ala Ala Ala Phe
 2595 2600

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 <213> Homo sapiens

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 35 40 45
 Ser Arg Gln Tyr Pro Val Phe Arg Gly Arg Pro Ser Gly Asn Glu Ser

50	55	60
Gln His Arg Leu Asp Phe Gln Leu Met Leu Lys Ile Arg Asp Thr Leu		
65	70	75
Tyr Ile Ala Gly Arg Asp Gln Val Tyr Thr Val Asn Leu Asn Glu Met		
85	90	95
Pro Lys Thr Glu Val Ile Pro Asn Lys Lys Leu Thr Trp Arg Ser Arg		
100	105	110
Gln Gln Asp Arg Glu Asn Cys Ala Met Lys Gly Lys His Lys Asp Glu		
115	120	125
Cys His Asn Phe Ile Lys Val Phe Val Pro Arg Asn Asp Glu Met Val		
130	135	140
Phe Val Cys Gly Thr Asn Ala Phe Asn Pro Met Cys Arg Tyr Tyr Arg		
145	150	155
Leu Ser Thr Leu Glu Tyr Asp Gly Glu Glu Ile Ser Gly Leu Ala Arg		
165	170	175
Cys Pro Phe Asp Ala Arg Gln Thr Asn Val Ala Leu Phe Ala Asp Gly		
180	185	190
Lys Leu Tyr Ser Ala Thr Val Ala Asp Phe Leu Ala Ser Asp Ala Val		
195	200	205
Ile Tyr Arg Ser Met Gly Asp Gly Ser Ala Leu Arg Thr Ile Lys Tyr		
210	215	220
Asp Ser Lys Trp Ile Lys Glu Pro His Phe Leu His Ala Ile Glu Tyr		
225	230	235
Gly Asn Tyr Val Tyr Phe Phe Arg Glu Ile Ala Val Glu His Asn		
245	250	255
Asn Leu Gly Lys Ala Val Tyr Ser Arg Val Ala Arg Ile Cys Lys Asn		
260	265	270
Asp Met Gly Gly Ser Gln Arg Val Leu Glu Lys His Trp Thr Ser Phe		
275	280	285
Leu Lys Ala Arg Leu Asn Cys Ser Val Pro Gly Asp Ser Phe Phe Tyr		
290	295	300
Phe Asp Val Leu Gln Ser Ile Thr Asp Ile Ile Gln Ile Asn Gly Ile		
305	310	315
Pro Thr Val Val Gly Val Phe Thr Thr Gln Leu Asn Ser Ile Pro Gly		
325	330	335
Ser Ala Val Cys Ala Phe Ser Met Asp Asp Ile Glu Lys Val Phe Lys		
340	345	350
Gly Arg Phe Lys Glu Gln Lys Thr Pro Asp Ser Val Trp Thr Ala Val		

355	360	365
Pro Glu Asp Lys Val Pro Lys Pro Arg Pro Gly Cys Cys Ala Lys His		
370	375	380
Gly Leu Ala Glu Ala Tyr Lys Thr Ser Ile Asp Phe Pro Asp Glu Thr		
385	390	395
Leu Ser Phe Ile Lys Ser His Pro Leu Met Asp Ser Ala Val Pro Pro		
405	410	415
Ile Ala Asp Glu Pro Trp Phe Thr Lys Thr Arg Val Arg Tyr Arg Leu		
420	425	430
Thr Ala Ile Ser Val Asp His Ser Ala Gly Pro Tyr Gln Asn Tyr Thr		
435	440	445
Val Ile Phe Val Gly Ser Glu Ala Gly Met Val Leu Lys Val Leu Ala		
450	455	460
Lys Thr Ser Pro Phe Ser Leu Asn Asp Ser Val Leu Leu Glu Glu Ile		
465	470	475
Glu Ala Tyr Asn His Ala Lys Cys Ser Ala Glu Asn Glu Glu Asp Lys		
485	490	495
Lys Val Ile Ser Leu Gln Leu Asp Lys Asp His His Ala Leu Tyr Val		
500	505	510
Ala Phe Ser Ser Cys Ile Ile Arg Ile Pro Leu Ser Arg Cys Glu Arg		
515	520	525
Tyr Gly Ser Cys Lys Lys Ser Cys Ile Ala Ser Arg Asp Pro Tyr Cys		
530	535	540
Gly Trp Leu Ser Gln Gly Ser Cys Gly Arg Val Thr Pro Gly Met Leu		
545	550	555
Leu Leu Thr Glu Asp Phe Ala Phe His Asn His Ser Ala Glu Gly		
565	570	575
Tyr Glu Gln Asp Thr Glu Phe Gly Asn Thr Ala His Leu Gly Asp Cys		
580	585	590
His Gly Val Arg Trp Glu Val Gln Ser Gly Glu Ser Asn Gln Met Val		
595	600	605
His Met Asn Val Leu Ile Thr Cys Val Phe Ala Ala Phe Val Leu Gly		
610	615	620
Ala Phe Ile Ala Gly Val Ala Val Tyr Cys Tyr Arg Asp Met Phe Val		
625	630	635
Arg Lys Asn Arg Lys Ile His Lys Asp Ala Glu Ser Ala Gln Ser Cys		
645	650	655
Thr Asp Ser Ser Gly Ser Phe Ala Lys Leu Asn Gly Leu Phe Asp Ser		

660	665	670
Pro Val Lys Glu Tyr Gln Gln Asn Ile Asp Ser Pro Lys Leu Tyr Ser		
675	680	685
Asn Leu Leu Thr Ser Arg Lys Glu Leu Pro Pro Asn Gly Asp Thr Lys		
690	695	700
Ser Met Val Met Asp His Arg Gly Gln Pro Pro Glu Leu Ala Ala Leu		
705	710	715
Pro Thr Pro Glu Ser Thr Pro Val Leu His Gln Lys Thr Leu Gln Ala		
725	730	735
Met Lys Ser His Ser Glu Lys Ala His Gly His Gly Ala Ser Arg Lys		
740	745	750
Glu Thr Pro Gln Phe Phe Pro Ser Ser Pro Pro Pro His Ser Pro Leu		
755	760	765
Ser His Gly His Ile Pro Ser Ala Ile Val Leu Pro Asn Ala Thr His		
770	775	780
Asp Tyr Asn Thr Ser Phe Ser Asn Ser Asn Ala His Lys Ala Glu Lys		
785	790	795
Lys Leu Gln Asn Ile Asp His Pro Leu Thr Lys Ser Ser Ser Lys Arg		
805	810	815
Asp His Arg Arg Ser Val Asp Ser Arg Asn Thr Leu Asn Asp Leu Leu		
820	825	830
Lys His Leu Asn Asp Pro Asn Ser Asn Pro Lys Ala Ile Met Gly Asp		
835	840	845
Ile Gln Met Ala His Gln Asn Leu Met Leu Asp Pro Met Gly Ser Met		
850	855	860
Ser Glu Val Pro Pro Lys Val Pro Asn Arg Glu Ala Ser Leu Tyr Ser		
865	870	875
Pro Pro Ser Thr Leu Pro Arg Asn Ser Pro Thr Lys Arg Val Asp Val		
885	890	895
Pro Thr Thr Pro Gly Val Pro Met Thr Ser Leu Glu Arg Gln Arg Gly		
900	905	910
Tyr His Lys Asn Ser Ser Gln Arg His Ser Ile Ser Ala Met Pro Lys		
915	920	925
Asn Leu Asn Ser Pro Asn Gly Val Leu Leu Ser Arg Gln Pro Ser Met		
930	935	940
Asn Arg Gly Gly Tyr Met Pro Thr Pro Thr Gly Ala Lys Val Asp Tyr		
945	950	955
Ile Gln Gly Thr Pro Val Ser Val His Leu Gln Pro Ser Leu Ser Arg		

965	970	975
Gln Ser Ser Tyr Thr Ser Asn Gly Thr Leu Pro Arg Thr Gly Leu Lys 980	985	990
Arg Thr Pro Ser Leu Lys Pro Asp Val Pro Pro Lys Pro Ser Phe Val 995	1000	1005
Pro Gln Thr Pro Ser Val Arg Pro Leu Asn Lys Tyr Thr Tyr 1010	1015	1020
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<213> Homo sapiens		
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Met Arg Val Phe Leu Leu Cys Ala Tyr Ile Leu Leu Met Val Ser 1	5	10
		15
Gln Leu Arg Ala Val Ser Phe Pro Glu Asp Asp Glu Pro Leu Asn Thr 20	25	30
Val Asp Tyr His Tyr Ser Arg Gln Tyr Pro Val Phe Arg Gly Arg Pro 35	40	45
Ser Gly Asn Glu Ser Gln His Arg Leu Asp Phe Gln Leu Met Leu Lys 50	55	60
Ile Arg Asp Thr Leu Tyr Ile Ala Gly Arg Asp Gln Val Tyr Thr Val 65	70	75
		80
Asn Leu Asn Glu Met Pro Lys Thr Glu Val Ile Pro Asn Lys Lys Leu 85	90	95
Thr Trp Arg Ser Arg Gln Gln Asp Arg Glu Asn Cys Ala Met Lys Gly 100	105	110
Lys His Lys Asp Glu Cys His Asn Phe Ile Lys Val Phe Val Pro Arg 115	120	125
Asn Asp Glu Met Val Phe Val Cys Gly Thr Asn Ala Phe Asn Pro Met 130	135	140
Cys Arg Tyr Tyr Arg Leu Ser Thr Leu Glu Tyr Asp Gly Glu Glu Ile 145	150	155
		160
Ser Gly Leu Ala Arg Cys Pro Phe Asp Ala Arg Gln Thr Asn Val Ala 165	170	175
Leu Phe Ala Asp Gly Lys Leu Tyr Ser Ala Thr Val Ala Asp Phe Leu 180	185	190
Ala Ser Asp Ala Val Ile Tyr Arg Ser Met Gly Asp Gly Ser Ala Leu 195	200	205

Arg Thr Ile Lys Tyr Asp Ser Lys Trp Ile Lys Glu Pro His Phe Leu
 210 215 220

His Ala Ile Glu Tyr Gly Asn Tyr Val Tyr Phe Phe Phe Arg Glu Ile
 225 230 235 240

Ala Val Glu His Asn Asn Leu Gly Lys Ala Val Tyr Ser Arg Val Ala
 245 250 255

Arg Ile Cys Lys Asn Asp Met Gly Gly Ser Gln Arg Val Leu Glu Lys
 260 265 270

His Trp Thr Ser Phe Leu Lys Ala Arg Leu Asn Cys Ser Val Pro Gly
 275 280 285

Asp Ser Phe Phe Tyr Phe Asp Val Leu Gln Ser Ile Thr Asp Ile Ile
 290 295 300

Gln Ile Asn Gly Ile Pro Thr Val Val Gly Val Phe Thr Thr Gln Leu
 305 310 315 320

Asn Ser Ile Pro Gly Ser Ala Val Cys Ala Phe Ser Met Asp Asp Ile
 325 330 335

Glu Lys Val Phe Lys Gly Arg Phe Lys Glu Gln Lys Thr Pro Asp Ser
 340 345 350

Val Trp Thr Ala Val Pro Glu Asp Lys Val Pro Lys Pro Arg Pro Gly
 355 360 365

Cys Cys Ala Lys His Gly Leu Ala Glu Ala Tyr Lys Thr Ser Ile Asp
 370 375 380

Phe Pro Asp Glu Thr Leu Ser Phe Ile Lys Ser His Pro Leu Met Asp
 385 390 395 400

Ser Ala Val Pro Pro Ile Ala Asp Glu Pro Trp Phe Thr Lys Thr Arg
 405 410 415

Val Arg Tyr Arg Leu Thr Ala Ile Ser Val Asp His Ser Ala Gly Pro
 420 425 430

Tyr Gln Asn Tyr Thr Val Ile Phe Val Gly Ser Glu Ala Gly Met Val
 435 440 445

Leu Lys Val Leu Ala Lys Thr Ser Pro Phe Ser Leu Asn Asp Ser Val
 450 455 460

Leu Leu Glu Glu Ile Glu Ala Tyr Asn His Ala Lys Cys Ser Ala Glu
 465 470 475 480

Asn Glu Glu Asp Lys Lys Val Ile Ser Leu Gln Leu Asp Lys Asp His
 485 490 495

His Ala Leu Tyr Val Ala Phe Ser Ser Cys Ile Ile Arg Ile Pro Leu
 500 505 510

Ser Arg Cys Glu Arg Tyr Gly Ser Cys Lys Lys Ser Cys Ile Ala Ser
 515 520 525
 Arg Asp Pro Tyr Cys Gly Trp Leu Ser Gln Gly Ser Cys Gly Arg Val
 530 535 540 540
 Thr Pro Gly Met Leu Leu Leu Thr Glu Asp Phe Phe Ala Phe His Asn
 545 550 555 560
 His Ser Ala Glu Gly Tyr Glu Gln Asp Thr Glu Phe Gly Asn Thr Ala
 565 570 575
 His Leu Gly Asp Cys His Gly Val Arg Trp Glu Val Gln Ser Gly Glu
 580 585 590
 Ser Asn Gln Met Val His Met Asn Val Leu Ile Thr Cys Val Phe Ala
 595 600 605
 Ala Phe Val Leu Gly Ala Phe Ile Ala Gly Val Ala Val Tyr Cys Tyr
 610 615 620
 Arg Asp Met Phe Val Arg Lys Asn Arg Lys Ile His Lys Asp Ala Glu
 625 630 635 640
 Ser Ala Gln Ser Cys Thr Asp Ser Ser Gly Ser Phe Ala Lys Leu Asn
 645 650 655
 Gly Leu Phe Asp Ser Pro Val Lys Glu Tyr Gln Gln Asn Ile Asp Ser
 660 665 670
 Pro Lys Leu Tyr Ser Asn Leu Leu Thr Ser Arg Lys Glu Leu Pro Pro
 675 680 685
 Asn Gly Asp Thr Lys Ser Met Val Met Asp His Arg Gly Gln Pro Pro
 690 695 700
 Glu Leu Ala Ala Leu Pro Thr Pro Glu Ser Thr Pro Val Leu His Gln
 705 710 715 720
 Lys Thr Leu Gln Ala Met Lys Ser His Ser Glu Lys Ala His Gly His
 725 730 735
 Gly Ala Ser Arg Lys Glu Thr Pro Gln Phe Phe Pro Ser Ser Pro Pro
 740 745 750
 Pro His Ser Pro Leu Ser His Gly His Ile Pro Ser Ala Ile Val Leu
 755 760 765
 Pro Asn Ala Thr His Asp Tyr Asn Thr Ser Phe Ser Asn Ser Asn Ala
 770 775 780
 His Lys Ala Glu Lys Lys Leu Gln Asn Ile Asp His Pro Leu Thr Lys
 785 790 795 800
 Ser Ser Ser Lys Arg Asp His Arg Arg Ser Val Asp Ser Arg Asn Thr
 805 810 815

Leu Asn Asp Leu Leu Lys His Leu Asn Asp Pro Asn Ser Asn Pro Lys
 820 825 830
 Ala Ile Met Gly Asp Ile Gln Met Ala His Gln Asn Leu Met Leu Asp
 835 840 845
 Pro Met Gly Ser Met Ser Glu Val Pro Pro Lys Val Pro Asn Arg Glu
 850 855 860
 Ala Ser Leu Tyr Ser Pro Pro Ser Thr Leu Pro Arg Asn Ser Pro Thr
 865 870 875 880
 Lys Arg Val Asp Val Pro Thr Thr Pro Gly Val Pro Met Thr Ser Leu
 885 890 895
 Glu Arg Gln Arg Gly Tyr His Lys Asn Ser Ser Gln Arg His Ser Ile
 900 905 910
 Ser Ala Met Pro Lys Asn Leu Asn Ser Pro Asn Gly Val Leu Leu Ser
 915 920 925
 Arg Gln Pro Ser Met Asn Arg Gly Gly Tyr Met Pro Thr Pro Thr Gly
 930 935 940
 Ala Lys Val Asp Tyr Ile Gln Gly Thr Pro Val Ser Val His Leu Gln
 945 950 955 960
 Pro Ser Leu Ser Arg Gln Ser Ser Tyr Thr Ser Asn Gly Thr Leu Pro
 965 970 975
 Arg Thr Gly Leu Lys Arg Thr Pro Ser Leu Lys Pro Asp Val Pro Pro
 980 985 990
 Lys Pro Ser Phe Val Pro Gln Thr Pro Ser Val Arg Pro Leu Asn Lys
 995 1000 1005
 Tyr Thr Tyr
 1010

<210> 62
 <211> 367
 <212> PRT
 <213> Homo sapiens

<400> 62
 Met Lys Gly Lys His Lys Asp Glu Cys His Asn Phe Ile Lys Val Phe
 1 5 10 15
 Val Pro Arg Asn Asp Glu Met Val Phe Val Cys Gly Thr Asn Ala Phe
 20 25 30
 Asn Pro Met Cys Arg Tyr Tyr Arg Leu Ser Thr Leu Glu Tyr Asp Gly
 35 40 45
 Glu Glu Ile Ser Gly Leu Ala Arg Cys Pro Phe Asp Ala Arg Gln Thr
 50 55 60

Asn Val Ala Leu Phe Ala Asp Gly Lys Leu Tyr Ser Ala Thr Val Ala
 65 70 75 80
 Asp Phe Leu Ala Ser Asp Ala Val Ile Tyr Arg Ser Met Gly Asp Gly
 85 90 95
 Ser Ala Leu Arg Thr Ile Lys Tyr Asp Ser Lys Trp Ile Lys Glu Pro
 100 105 110
 His Phe Leu His Ala Ile Glu Tyr Gly Asn Tyr Val Tyr Phe Phe Phe
 115 120 125
 Arg Glu Ile Ala Val Glu His Asn Asn Leu Gly Lys Ala Val Tyr Ser
 130 135 140
 Arg Val Ala Arg Ile Cys Lys Asn Asp Met Gly Gly Ser Gln Arg Val
 145 150 155 160
 Leu Glu Lys His Trp Thr Ser Phe Leu Lys Ala Arg Leu Asn Cys Ser
 165 170 175
 Val Pro Gly Asp Ser Phe Phe Tyr Phe Asp Val Leu Gln Ser Ile Thr
 180 185 190
 Asp Ile Ile Gln Ile Asn Gly Ile Pro Thr Val Val Gly Val Phe Thr
 195 200 205
 Thr Gln Leu Asn Ser Ile Pro Gly Ser Ala Val Cys Ala Phe Ser Met
 210 215 220
 Asp Asp Ile Glu Lys Val Phe Lys Gly Arg Phe Lys Glu Gln Lys Thr
 225 230 235 240
 Pro Asp Ser Val Trp Thr Ala Val Pro Glu Asp Lys Val Pro Lys Pro
 245 250 255
 Arg Pro Gly Cys Cys Ala Lys His Gly Leu Ala Glu Ala Tyr Lys Thr
 260 265 270
 Ser Ile Asp Phe Pro Asp Glu Thr Leu Ser Phe Ile Lys Ser His Pro
 275 280 285
 Leu Met Asp Ser Ala Val Pro Pro Ile Ala Asp Glu Pro Trp Phe Thr
 290 295 300
 Lys Thr Arg Val Arg Tyr Arg Leu Thr Ala Ile Ser Val Asp His Ser
 305 310 315 320
 Ala Gly Pro Tyr Gln Asn Tyr Thr Val Ile Phe Val Gly Ser Glu Ala
 325 330 335
 Gly Met Val Leu Lys Val Leu Ala Lys Thr Ser Pro Phe Ser Leu Asn
 340 345 350
 Asp Ser Val Leu Leu Glu Glu Ile Glu Ala Tyr Asn His Ala Lys
 355 360 365

<210> 63
<211> 1030
<212> PRT
<213> Homo sapiens

<400> 63
Met Arg Ser Glu Ala Leu Leu Leu Tyr Phe Thr Leu Leu His Phe Ala
1 5 10 15
Gly Ala Gly Phe Pro Glu Asp Ser Glu Pro Ile Ser Ile Ser His Gly
20 25 30
Asn Tyr Thr Lys Gln Tyr Pro Val Phe Val Gly His Lys Pro Gly Arg
35 40 45
Asn Thr Thr Gln Arg His Arg Leu Asp Ile Gln Met Ile Met Ile Met
50 55 60
Asn Gly Thr Leu Tyr Ile Ala Ala Arg Asp His Ile Tyr Thr Val Asp
65 70 75 80
Ile Asp Thr Ser His Thr Glu Glu Ile Tyr Cys Ser Lys Lys Leu Thr
85 90 95
Trp Lys Ser Arg Gln Ala Asp Val Asp Thr Cys Arg Met Lys Gly Lys
100 105 110
His Lys Asp Glu Cys His Asn Phe Ile Lys Val Leu Leu Lys Lys Asn
115 120 125
Asp Asp Ala Leu Phe Val Cys Gly Thr Asn Ala Phe Asn Pro Ser Cys
130 135 140
Arg Asn Tyr Lys Met Asp Thr Leu Glu Pro Phe Gly Asp Glu Phe Ser
145 150 155 160
Gly Met Ala Arg Cys Pro Tyr Asp Ala Lys His Ala Asn Val Ala Leu
165 170 175
Phe Ala Asp Gly Lys Leu Tyr Ser Ala Thr Val Thr Asp Phe Leu Ala
180 185 190
Ile Asp Ala Val Ile Tyr Arg Ser Leu Gly Glu Ser Pro Thr Leu Arg
195 200 205
Thr Val Lys His Asp Ser Lys Trp Leu Lys Glu Pro Tyr Phe Val Gln
210 215 220
Ala Val Asp Tyr Gly Asp Tyr Ile Tyr Phe Phe Arg Glu Ile Ala
225 230 235 240
Val Glu Tyr Asn Thr Met Gly Lys Val Val Phe Pro Arg Val Ala Gln
245 250 255
Val Cys Lys Asn Asp Met Gly Gly Ser Gln Arg Val Leu Glu Lys Gln

	260	265	270
Trp Thr Ser Phe Leu Lys Ala Arg Leu Asn Cys Ser Val Pro Gly Asp			
275	280	285	
Ser His Phe Tyr Phe Asn Ile Leu Gln Ala Val Thr Asp Val Ile Arg			
290	295	300	
Ile Asn Gly Arg Asp Val Val Leu Ala Thr Phe Ser Thr Pro Tyr Asn			
305	310	315	320
Ser Ile Pro Gly Ser Ala Val Cys Ala Tyr Asp Met Leu Asp Ile Ala			
325	330	335	
Ser Val Phe Thr Gly Arg Phe Lys Glu Gln Lys Ser Pro Asp Ser Thr			
340	345	350	
Trp Thr Pro Val Pro Asp Glu Arg Val Pro Lys Pro Arg Pro Gly Cys			
355	360	365	
Cys Ala Gly Ser Ser Ser Leu Glu Arg Tyr Ala Thr Ser Asn Glu Phe			
370	375	380	
Pro Asp Asp Thr Leu Asn Phe Ile Lys Thr His Pro Leu Met Asp Glu			
385	390	395	400
Ala Val Pro Ser Ile Phe Asn Arg Pro Trp Phe Leu Arg Thr Met Val			
405	410	415	
Arg Tyr Arg Leu Thr Lys Ile Ala Val Asp Thr Ala Ala Gly Pro Tyr			
420	425	430	
Gln Asn His Thr Val Val Phe Leu Gly Ser Glu Lys Gly Ile Ile Leu			
435	440	445	
Lys Phe Leu Ala Arg Ile Gly Asn Ser Gly Phe Leu Asn Asp Ser Leu			
450	455	460	
Phe Leu Glu Glu Met Ser Val Tyr Asn Ser Glu Lys Cys Ser Tyr Asp			
465	470	475	480
Gly Val Glu Asp Lys Arg Ile Met Gly Met Gln Leu Asp Arg Ala Ser			
485	490	495	
Ser Ser Leu Tyr Val Ala Phe Ser Thr Cys Val Ile Lys Val Pro Leu			
500	505	510	
Gly Arg Cys Glu Arg His Gly Lys Cys Lys Lys Thr Cys Ile Ala Ser			
515	520	525	
Arg Asp Pro Tyr Cys Gly Trp Ile Lys Glu Gly Ala Cys Ser His			
530	535	540	
Leu Ser Pro Asn Ser Arg Leu Thr Phe Glu Gln Asp Ile Glu Arg Gly			
545	550	555	560
Asn Thr Asp Gly Leu Gly Asp Cys His Asn Ser Phe Val Ala Leu Asn			

565	570	575
Gly His Ser Ser Ser Leu Leu Pro Ser Thr Thr Thr Ser Asp Ser Thr		
580	585	590
Ala Gln Glu Gly Tyr Glu Ser Arg Gly Gly Met Leu Asp Trp Lys His		
595	600	605
Leu Leu Asp Ser Pro Asp Ser Thr Asp Pro Leu Gly Ala Val Ser Ser		
610	615	620
His Asn His Gln Asp Lys Lys Gly Val Ile Arg Glu Ser Tyr Leu Lys		
625	630	635
Gly His Asp Gln Leu Val Pro Val Thr Leu Leu Ala Ile Ala Val Ile		
645	650	655
Leu Ala Phe Val Met Gly Ala Val Phe Ser Gly Ile Thr Val Tyr Cys		
660	665	670
Val Cys Asp His Arg Arg Lys Asp Val Ala Val Val Gln Arg Lys Glu		
675	680	685
Lys Glu Leu Thr His Ser Arg Arg Gly Ser Met Ser Ser Val Thr Lys		
690	695	700
Leu Ser Gly Leu Phe Gly Asp Thr Gln Ser Lys Asp Pro Lys Pro Glu		
705	710	720
Ala Ile Leu Thr Pro Leu Met His Asn Gly Lys Leu Ala Thr Pro Gly		
725	730	735
Asn Thr Ala Lys Met Leu Ile Lys Ala Asp Gln His His Leu Asp Leu		
740	745	750
Thr Ala Leu Pro Thr Pro Glu Ser Thr Pro Thr Leu Gln Gln Lys Arg		
755	760	765
Lys Pro Ser Arg Gly Ser Arg Glu Trp Glu Arg Asn Gln Asn Leu Ile		
770	775	780
Asn Ala Cys Thr Lys Asp Met Pro Pro Met Gly Ser Pro Val Ile Pro		
785	790	800
Thr Asp Leu Pro Leu Arg Ala Ser Pro Ser His Ile Pro Ser Val Val		
805	810	815
Val Leu Pro Ile Thr Gln Gln Gly Tyr Gln His Glu Tyr Val Asp Gln		
820	825	830
Pro Lys Met Ser Glu Val Ala Gln Met Ala Leu Glu Asp Gln Ala Ala		
835	840	845
Thr Leu Glu Tyr Lys Thr Ile Lys Glu His Leu Ser Ser Lys Ser Pro		
850	855	860
Asn His Gly Val Asn Leu Val Glu Asn Leu Asp Ser Leu Pro Pro Lys		

865	870	875	880
Val Pro Gln Arg Glu Ala Ser Leu Gly Pro Pro Gly Ala Ser Leu Ser			
885		890	895
Gln Thr Gly Leu Ser Lys Arg Leu Glu Met His His Ser Ser Ser Tyr			
900		905	910
Gly Val Asp Tyr Lys Arg Ser Tyr Pro Thr Asn Ser Leu Thr Arg Ser			
915		920	925
His Gln Ala Thr Thr Leu Lys Arg Asn Asn Thr Asn Ser Ser Asn Ser			
930		935	940
Ser His Leu Ser Arg Asn Gln Ser Phe Gly Arg Gly Asp Asn Pro Pro			
945		950	955
Pro Ala Pro Gln Arg Val Asp Ser Ile Gln Val His Ser Ser Gln Pro			
965		970	975
Ser Gly Gln Ala Val Thr Val Ser Arg Gln Pro Ser Leu Asn Ala Tyr			
980		985	990
Asn Ser Leu Thr Arg Ser Gly Leu Lys Arg Thr Pro Ser Leu Lys Pro			
995		1000	1005
Asp Val Pro Pro Lys Pro Ser Phe Ala Pro Leu Ser Thr Ser Met Lys			
1010		1015	1020
Pro Asn Asp Ala Cys Thr			
1025		1030	
<210> 64			
<211> 888			
<212> PRT			
<213> Homo sapiens			
<400> 64			
Met Arg Pro Ala Ala Leu Leu Leu Cys Leu Thr Leu Leu His Cys Ala			
1	5	10	15
Gly Ala Gly Phe Pro Glu Asp Ser Glu Pro Ile Ser Ile Ser His Gly			
20		25	30
Asn Tyr Thr Lys Gln Tyr Pro Val Phe Val Gly His Lys Pro Gly Arg			
35		40	45
Asn Thr Thr Gln Arg His Arg Leu Asp Ile Gln Met Ile Met Ile Met			
50		55	60
Asn Arg Thr Leu Tyr Val Ala Ala Arg Asp His Ile Tyr Thr Val Asp			
65		70	75
Ile Asp Thr Ser His Thr Glu Glu Ile Tyr Cys Ser Lys Lys Leu Thr			
85		90	95

Trp Lys Ser Arg Gln Ala Asp Val Asp Thr Cys Arg Met Lys Gly Lys
 100 105 110
 His Lys Asp Glu Cys His Asn Phe Ile Lys Val Leu Leu Lys Lys Asn
 115 120 125
 Asp Asp Thr Leu Phe Val Cys Gly Thr Asn Ala Phe Asn Pro Ser Cys
 130 135 140
 Arg Asn Tyr Arg Val Asp Thr Leu Glu Thr Phe Gly Asp Glu Phe Ser
 145 150 155 160
 Gly Met Ala Arg Cys Pro Tyr Asp Ala Lys His Ala Asn Ile Ala Leu
 165 170 175
 Phe Ala Asp Gly Lys Leu Tyr Ser Ala Thr Val Thr Asp Phe Leu Ala
 180 185 190
 Ile Asp Ala Val Ile Tyr Arg Ser Leu Gly Asp Ser Pro Thr Leu Arg
 195 200 205
 Thr Val Lys His Asp Ser Lys Trp Leu Lys Glu Pro Tyr Phe Val Gln
 210 215 220
 Ala Val Asp Tyr Gly Asp Tyr Ile Tyr Phe Phe Arg Glu Ile Ala
 225 230 235 240
 Val Glu Tyr Asn Thr Met Gly Lys Val Val Phe Pro Arg Val Ala Gln
 245 250 255
 Val Cys Lys Asn Asp Met Gly Gly Ser Gln Arg Val Leu Glu Lys Gln
 260 265 270
 Trp Thr Ser Phe Leu Lys Ala Arg Leu Asn Cys Ser Val Pro Gly Asp
 275 280 285
 Ser His Phe Tyr Phe Asn Ile Leu Gln Ala Val Thr Asp Val Ile Arg
 290 295 300
 Ile Asn Gly Arg Asp Val Val Leu Ala Thr Phe Ser Thr Pro Tyr Asn
 305 310 315 320
 Ser Ile Pro Gly Ser Ala Val Cys Ala Tyr Asp Met Leu Asp Ile Ala
 325 330 335
 Asn Val Phe Thr Gly Arg Phe Lys Glu Gln Lys Ser Pro Asp Ser Thr
 340 345 350
 Trp Thr Pro Val Pro Asp Glu Arg Val Pro Lys Pro Arg Pro Gly Cys
 355 360 365
 Cys Ala Gly Ser Ser Ser Leu Glu Lys Tyr Ala Thr Ser Asn Glu Phe
 370 375 380
 Pro Asp Asp Thr Leu Asn Phe Ile Lys Thr His Pro Leu Met Asp Glu
 385 390 395 400

Ala Val Pro Ser Ile Ile Asn Arg Pro Trp Phe Leu Arg Thr Met Val
405 410 415

Arg Tyr Arg Leu Thr Lys Ile Ala Val Asp Asn Ala Ala Gly Pro Tyr
420 425 430

Gln Asn His Thr Val Val Phe Leu Gly Ser Glu Lys Gly Ile Ile Leu
435 440 445

Lys Phe Leu Ala Arg Ile Gly Ser Ser Gly Phe Leu Asn Gly Ser Leu
450 455 460

Phe Leu Glu Glu Met Asn Val Tyr Asn Pro Glu Lys Cys Ser Tyr Asp
465 470 475 480

Gly Val Glu Asp Lys Arg Ile Met Gly Met Gln Leu Asp Arg Ala Ser
485 490 495

Gly Ser Leu Tyr Val Ala Phe Ser Thr Cys Val Ile Lys Val Pro Leu
500 505 510

Gly Arg Cys Glu Arg His Gly Lys Cys Lys Lys Thr Cys Ile Ala Ser
515 520 525

Arg Asp Pro Tyr Cys Gly Trp Val Arg Glu Ser Gly Ser Cys Ala His
530 535 540

Leu Ser Pro Leu Ser Arg Leu Thr Phe Glu Gln Asp Ile Glu Arg Gly
545 550 555 560

Asn Thr Asp Gly Leu Gly Asp Cys His Asn Ser Phe Val Ala Leu Asn
565 570 575

Gly His Ala Ser Ser Leu Tyr Pro Ser Thr Thr Ser Asp Ser Ala
580 585 590

Ser Arg Asp Gly Tyr Glu Ser Arg Gly Gly Met Leu Asp Trp Asn Asp
595 600 605

Leu Leu Glu Ala Pro Gly Ser Thr Asp Pro Leu Gly Ala Val Ser Ser
610 615 620

His Asn His Gln Asp Lys Lys Gly Val Ile Arg Glu Ser Tyr Leu Lys
625 630 635 640

Ser Asn Asp Gln Leu Val Pro Val Thr Leu Leu Ala Ile Ala Val Ile
645 650 655

Leu Ala Phe Val Met Gly Ala Val Phe Ser Gly Ile Ile Val Tyr Cys
660 665 670

Val Cys Asp His Arg Arg Lys Asp Val Ala Val Val Gln Arg Lys Glu
675 680 685

Lys Glu Leu Thr His Ser Arg Arg Gly Ser Met Ser Ser Val Thr Lys
690 695 700

Leu Ser Gly Leu Phe Gly Asp Thr Gln Ser Lys Asp Pro Lys Pro Glu
 705 710 715 720
 Ala Ile Leu Thr Pro Leu Met His Asn Gly Lys Leu Ala Thr Pro Ser
 725 730 735
 Asn Thr Ala Lys Met Leu Ile Lys Ala Asp Gln His His Leu Asp Leu
 740 745 750
 Thr Ala Leu Pro Thr Pro Glu Ser Thr Pro Thr Leu Gln Glu Lys Arg
 755 760 765
 Lys Pro Asn Arg Gly Ser Arg Glu Trp Glu Arg Asn Gln Asn Ile Ile
 770 775 780
 Asn Ala Cys Thr Lys Asp Met Pro Pro Met Gly Ser Pro Val Ile Pro
 785 790 795 800
 Thr Asp Leu Pro Leu Arg Ala Ser Pro Ser His Ile Pro Ser Val Val
 805 810 815
 Val Leu Pro Ile Thr Gln Gln Gly Tyr Gln His Glu Tyr Val Asp Gln
 820 825 830
 Pro Lys Met Ser Glu Val Val Ala Gln Met Ala Leu Glu Asp Gln Ala
 835 840 845
 Ala Thr Leu Glu Tyr Lys Thr Ile Lys Glu His Leu Ser Ser Glu Ser
 850 855 860
 Ser Pro Tyr Val Leu Lys Gln Phe Ser Glu Ala Phe Asn Arg Gln Gly
 865 870 875 880
 Ile Ile Leu Ser Val Ala Val Glu
 885

<210> 65
 <211> 30
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence:
 oligonucleotide primer

 <400> 65

gatctcccgaaaccctctg agccgaaggg 30

<210> 66
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence:

oligonucleotide primer

<400> 66
ggcagcgccc tacacggt 18

<210> 67
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
oligonucleotide primer

<400> 67
gatgagtgcg cgactggc 18

<210> 68
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
oligonucleotide primer

<400> 68
cctcagcgtc cgcctcct 18

<210> 69
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
oligonucleotide primer

<400> 69
cgcaactcatc cacatcttcg c 21

<210> 70
<211> 703
<212> PRT
<213> Homo sapiens

<400> 70
Pro Glu Glu Gln Pro His Val Gly Asn Tyr Arg Leu Leu Arg Thr Ile
1 5 10 15
Gly Lys Gly Asn Phe Ala Lys Val Lys Leu Ala Arg His Ile Leu Thr
20 25 30

Gly Arg Glu Val Ala Ile Lys Ile Ile Asp Lys Thr Gln Leu Asn Pro
 35 40 45

 Ser Ser Leu Gln Lys Leu Phe Arg Glu Val Arg Ile Met Lys Gly Leu
 50 55 60

 Asn His Pro Asn Ile Val Lys Leu Phe Glu Val Ile Glu Thr Glu Lys
 65 70 75 80

 Thr Leu Tyr Leu Val Met Glu Tyr Ala Ser Ala Gly Glu Val Phe Asp
 85 90 95

 Tyr Leu Val Ser His Gly Arg Met Lys Glu Lys Glu Ala Arg Ala Lys
 100 105 110

 Phe Arg Gln Ile Val Ser Ala Val His Tyr Cys His Gln Lys Asn Ile
 115 120 125

 Val His Arg Asp Leu Lys Ala Glu Asn Leu Leu Asp Ala Glu Ala
 130 135 140

 Asn Ile Lys Ile Ala Asp Phe Gly Phe Ser Asn Glu Phe Thr Leu Gly
 145 150 155 160

 Ser Lys Leu Asp Thr Phe Cys Gly Ser Pro Pro Tyr Ala Ala Pro Glu
 165 170 175

 Leu Phe Gln Gly Lys Lys Tyr Asp Gly Pro Glu Val Asp Ile Trp Ser
 180 185 190

 Leu Gly Val Ile Leu Tyr Thr Leu Val Ser Gly Ser Leu Pro Phe Asp
 195 200 205

 Gly His Asn Leu Lys Glu Leu Arg Glu Arg Val Leu Arg Gly Lys Tyr
 210 215 220

 Arg Val Pro Phe Tyr Met Ser Thr Asp Cys Glu Ser Ile Leu Arg Arg
 225 230 235 240

 Phe Leu Val Leu Asn Pro Ala Lys Arg Cys Thr Leu Glu Gln Ile Met
 245 250 255

 Lys Asp Lys Trp Ile Asn Ile Gly Tyr Glu Gly Glu Glu Leu Lys Pro
 260 265 270

 Tyr Thr Glu Pro Glu Glu Asp Phe Gly Asp Thr Lys Arg Ile Glu Val
 275 280 285

 Met Val Gly Met Gly Tyr Thr Arg Glu Glu Ile Lys Glu Ser Leu Thr
 290 295 300

 Ser Gln Lys Tyr Asn Glu Val Thr Ala Thr Tyr Leu Leu Leu Gly Arg
 305 310 315 320

 Lys Thr Glu Glu Gly Gly Asp Arg Gly Ala Pro Gly Leu Ala Leu Ala
 325 330 335

Arg Val Arg Ala Pro Ser Asp Thr Thr Asn Gly Thr Ser Ser Ser Lys
 340 345 350

 Gly Thr Ser His Ser Lys Gly Gln Arg Ser Ser Ser Thr Tyr His
 355 360 365

 Arg Gln Arg Arg His Ser Asp Phe Cys Gly Pro Ser Pro Ala Pro Leu
 370 375 380

 His Pro Lys Arg Ser Pro Thr Ser Thr Gly Glu Ala Glu Leu Lys Glu
 385 390 395 400

 Glu Arg Leu Pro Gly Arg Lys Ala Ser Cys Ser Thr Ala Gly Ser Gly
 405 410 415

 Ser Arg Gly Leu Pro Pro Ser Ser Pro Met Val Ser Ser Ala His Asn
 420 425 430

 Pro Asn Lys Ala Glu Ile Pro Glu Arg Arg Lys Asp Ser Thr Ser Thr
 435 440 445

 Pro Asn Asn Leu Pro Pro Ser Met Met Thr Arg Arg Asn Thr Tyr Val
 450 455 460

 Cys Thr Glu Arg Pro Gly Ala Glu Arg Pro Ser Leu Leu Pro Asn Gly
 465 470 475 480

 Lys Glu Asn Ser Ser Gly Thr Pro Arg Val Pro Pro Ala Ser Pro Ser
 485 490 495

 Ser His Ser Leu Ala Pro Pro Ser Gly Glu Arg Ser Arg Leu Ala Arg
 500 505 510

 Gly Ser Thr Ile Arg Ser Thr Phe His Gly Gly Gln Val Arg Asp Arg
 515 520 525

 Arg Ala Gly Gly Gly Gly Gly Val Gln Asn Gly Pro Pro Ala
 530 535 540

 Ser Pro Thr Leu Ala His Glu Ala Ala Pro Leu Pro Ala Gly Arg Pro
 545 550 555 560

 Arg Pro Thr Thr Asn Leu Phe Thr Lys Leu Thr Ser Lys Leu Thr Arg
 565 570 575

 Arg Val Ala Asp Glu Pro Glu Arg Ile Gly Gly Pro Glu Val Thr Ser
 580 585 590

 Cys His Leu Pro Trp Asp Gln Thr Glu Thr Ala Pro Arg Leu Leu Arg
 595 600 605

 Phe Pro Trp Ser Val Lys Leu Thr Ser Ser Arg Pro Pro Glu Ala Leu
 610 615 620

 Met Ala Ala Leu Arg Gln Ala Thr Ala Ala Ala Arg Cys Arg Cys Arg
 625 630 635 640

Gln Pro Gln Pro Phe Leu Leu Ala Cys Leu His Gly Gly Ala Gly Gly
645 650 655

Pro Glu Pro Leu Ser His Phe Glu Val Glu Val Cys Gln Leu Pro Arg
660 665 670

Pro Gly Leu Arg Gly Val Leu Phe Arg Arg Val Ala Gly Thr Ala Leu
675 680 685

Ala Phe Arg Thr Leu Val Thr Arg Ile Ser Asn Asp Leu Glu Leu
690 695 700

<210> 71

<211> 639

<212> PRT

<213> Homo sapiens

<400> 71
Pro Glu Glu Gln Pro His Val Gly Asn Tyr Arg Leu Leu Arg Thr Ile
1 5 10 15

Gly Lys Gly Asn Phe Ala Lys Val Lys Leu Ala Arg His Ile Leu Thr
20 25 30

Gly Arg Glu Val Ala Ile Lys Ile Ile Asp Lys Thr Gln Leu Asn Pro
35 40 45

Ser Ser Leu Gln Lys Leu Phe Arg Glu Val Arg Ile Met Lys Gly Leu
50 55 60

Asn His Pro Asn Ile Val Lys Leu Phe Glu Val Ile Glu Thr Glu Lys
65 70 75 80

Thr Leu Tyr Leu Val Met Glu Tyr Ala Ser Ala Gly Glu Val Phe Asp
85 90 95

Tyr Leu Val Ser His Gly Arg Met Lys Glu Lys Glu Ala Arg Ala Lys
100 105 110

Phe Arg Gln Ile Val Ser Ala Val His Tyr Cys His Gln Lys Asn Ile
115 120 125

Val His Arg Asp Leu Lys Ala Glu Asn Leu Leu Leu Asp Ala Glu Ala
130 135 140

Asn Ile Lys Ile Ala Asp Phe Gly Phe Ser Asn Glu Phe Thr Leu Gly
145 150 155 160

Ser Lys Leu Asp Thr Phe Cys Gly Ser Pro Pro Tyr Ala Ala Pro Glu
165 170 175

Leu Phe Gln Gly Lys Lys Tyr Asp Gly Pro Glu Val Asp Ile Trp Ser
180 185 190

Leu Gly Val Ile Leu Tyr Thr Leu Val Ser Gly Ser Leu Pro Phe Asp
195 200 205

Gly His Asn Leu Lys Glu Leu Arg Glu Arg Val Leu Arg Gly Lys Tyr
 210 215 220
 Arg Val Pro Phe Tyr Met Ser Thr Asp Cys Glu Ser Ile Leu Arg Arg
 225 230 235 240
 Phe Leu Val Leu Asn Pro Ala Lys Arg Cys Thr Leu Glu Gln Ile Met
 245 250 255
 Lys Asp Lys Trp Ile Asn Ile Gly Tyr Glu Gly Glu Glu Leu Lys Pro
 260 265 270
 Tyr Thr Glu Pro Glu Glu Asp Phe Gly Asp Thr Lys Arg Ile Glu Val
 275 280 285
 Met Val Gly Met Gly Tyr Thr Arg Glu Glu Ile Lys Glu Ser Leu Thr
 290 295 300
 Ser Gln Lys Tyr Asn Glu Val Thr Ala Thr Tyr Leu Leu Leu Gly Arg
 305 310 315 320
 Lys Thr Glu Glu Gly Asp Arg Gly Ala Pro Gly Leu Ala Leu Ala
 325 330 335
 Arg Val Arg Ala Pro Ser Asp Thr Thr Asn Gly Thr Ser Ser Ser Lys
 340 345 350
 Gly Thr Ser His Ser Lys Gly Gln Arg Ser Ser Ser Ser Thr Tyr His
 355 360 365
 Arg Gln Arg Arg His Ser Asp Phe Cys Gly Pro Ser Pro Ala Pro Leu
 370 375 380
 His Pro Lys Arg Ser Pro Thr Ser Thr Gly Glu Ala Glu Leu Lys Glu
 385 390 395 400
 Glu Arg Leu Pro Gly Arg Lys Ala Ser Cys Ser Thr Ala Gly Ser Gly
 405 410 415
 Ser Arg Gly Leu Pro Pro Ser Ser Pro Met Val Ser Ser Ala His Asn
 420 425 430
 Pro Asn Lys Ala Glu Ile Pro Glu Arg Arg Lys Asp Ser Thr Ser Thr
 435 440 445
 Pro Asn Asn Leu Pro Pro Ser Met Met Thr Arg Arg Asn Thr Tyr Val
 450 455 460
 Cys Thr Glu Arg Pro Gly Ala Glu Arg Pro Ser Leu Leu Pro Asn Gly
 465 470 475 480
 Lys Glu Asn Ser Ser Gly Thr Pro Arg Val Pro Pro Ala Ser Pro Ser
 485 490 495
 Ser His Ser Leu Ala Pro Pro Ser Gly Glu Arg Ser Arg Leu Ala Arg
 500 505 510

Gly Ser Thr Ile Arg Ser Thr Phe His Gly Gly Gln Val Arg Asp Arg
 515 520 525
 Arg Ala Gly Gly Gly Gly Gly Val Gln Asn Gly Pro Pro Ala
 530 535 540
 Ser Pro Thr Leu Ala His Glu Ala Ala Pro Leu Pro Ala Gly Arg Pro
 545 550 555 560
 Arg Pro Thr Thr Asn Leu Phe Thr Lys Leu Thr Ser Lys Leu Thr Arg
 565 570 575
 Arg Val Thr Leu Asp Pro Ser Lys Arg Gln Asn Ser Asn Arg Cys Val
 580 585 590
 Ser Gly Ala Ser Leu Pro Gln Gly Ser Lys Ile Arg Ser Gln Thr Asn
 595 600 605
 Leu Arg Glu Ser Gly Asp Leu Arg Ser Gln Val Ala Ile Tyr Leu Gly
 610 615 620
 Ile Lys Arg Lys Pro Pro Pro Gly Cys Ser Asp Ser Pro Gly Val
 625 630 635

<210> 72
 <211> 639
 <212> PRT
 <213> Homo sapiens

<400> 72
 Pro Glu Glu Gln Pro His Val Gly Asn Tyr Arg Leu Leu Arg Thr Ile
 1 5 10 15
 Gly Lys Gly Asn Phe Ala Lys Val Lys Leu Ala Arg His Ile Leu Thr
 20 25 30
 Gly Arg Glu Val Ala Ile Lys Ile Ile Asp Lys Thr Gln Leu Asn Pro
 35 40 45
 Ser Ser Leu Gln Lys Leu Phe Arg Glu Val Arg Ile Met Lys Gly Leu
 50 55 60
 Asn His Pro Asn Ile Val Lys Leu Phe Glu Val Ile Glu Thr Glu Lys
 65 70 75 80
 Thr Leu Tyr Leu Val Met Glu Tyr Ala Ser Ala Gly Glu Val Phe Asp
 85 90 95
 Tyr Leu Val Ser His Gly Arg Met Lys Glu Lys Glu Ala Arg Ala Lys
 100 105 110
 Phe Arg Gln Ile Val Ser Ala Val His Tyr Cys His Gln Lys Asn Ile
 115 120 125
 Val His Arg Asp Leu Lys Ala Glu Asn Leu Leu Leu Asp Ala Glu Ala

130	135	140
Asn Ile Lys Ile Ala Asp Phe Gly Phe Ser Asn Glu Phe Thr Leu Gly		
145	150	155
Ser Lys Leu Asp Thr Phe Cys Gly Ser Pro Pro Tyr Ala Ala Pro Glu		
165	170	175
Leu Phe Gln Gly Lys Lys Tyr Asp Gly Pro Glu Val Asp Ile Trp Ser		
180	185	190
Leu Gly Val Ile Leu Tyr Thr Leu Val Ser Gly Ser Leu Pro Phe Asp		
195	200	205
Gly His Asn Leu Lys Glu Leu Arg Glu Arg Val Leu Arg Gly Lys Tyr		
210	215	220
Arg Val Pro Phe Tyr Met Ser Thr Asp Cys Glu Ser Ile Leu Arg Arg		
225	230	235
Phe Leu Val Leu Asn Pro Ala Lys Arg Cys Thr Leu Glu Gln Ile Met		
245	250	255
Lys Asp Lys Trp Ile Asn Ile Gly Tyr Glu Gly Glu Glu Leu Lys Pro		
260	265	270
Tyr Thr Glu Pro Glu Glu Asp Phe Gly Asp Thr Lys Arg Ile Glu Val		
275	280	285
Met Val Gly Met Gly Tyr Thr Arg Glu Glu Ile Lys Glu Ser Leu Thr		
290	295	300
Ser Gln Lys Tyr Asn Glu Val Thr Ala Thr Tyr Leu Leu Leu Gly Arg		
305	310	315
Lys Thr Glu Glu Gly Gly Asp Arg Gly Ala Pro Gly Leu Ala Leu Ala		
325	330	335
Arg Val Arg Ala Pro Ser Asp Thr Thr Asn Gly Thr Ser Ser Ser Lys		
340	345	350
Gly Thr Ser His Ser Lys Gly Gln Arg Ser Ser Ser Thr Tyr His		
355	360	365
Arg Gln Arg Arg His Ser Asp Phe Cys Gly Pro Ser Pro Ala Pro Leu		
370	375	380
His Pro Lys Arg Ser Pro Thr Ser Thr Gly Glu Ala Glu Leu Lys Glu		
385	390	395
Glu Arg Leu Pro Gly Arg Lys Ala Ser Cys Ser Thr Ala Gly Ser Gly		
405	410	415
Ser Arg Gly Leu Pro Pro Ser Ser Pro Met Val Ser Ser Ala His Asn		
420	425	430
Pro Asn Lys Ala Glu Ile Pro Glu Arg Arg Lys Asp Ser Thr Ser Thr		

435	440	445
Pro Asn Asn Leu Pro Pro Ser Met Met Thr Arg Arg Asn Thr Tyr Val		
450	455	460
Cys Thr Glu Arg Pro Gly Ala Glu Arg Pro Ser Leu Leu Pro Asn Gly		
465	470	475
Lys Glu Asn Ser Ser Gly Thr Pro Arg Val Pro Pro Ala Ser Pro Ser		
485	490	495
Ser His Ser Leu Ala Pro Pro Ser Gly Glu Arg Ser Arg Leu Ala Arg		
500	505	510
Gly Ser Thr Ile Arg Ser Thr Phe His Gly Gly Gln Val Arg Asp Arg		
515	520	525
Arg Ala Gly Gly Gly Gly Gly Val Gln Asn Gly Pro Pro Ala		
530	535	540
Ser Pro Thr Leu Ala His Glu Ala Ala Pro Leu Pro Ala Gly Arg Pro		
545	550	555
Arg Pro Thr Thr Asn Leu Phe Thr Lys Leu Thr Ser Lys Leu Thr Arg		
565	570	575
Arg Val Thr Leu Asp Pro Ser Lys Arg Gln Asn Ser Asn Arg Cys Val		
580	585	590
Ser Gly Ala Ser Leu Pro Gln Gly Ser Lys Ile Arg Ser Gln Thr Asn		
595	600	605
Leu Arg Glu Ser Gly Asp Leu Arg Ser Gln Val Ala Ile Tyr Leu Gly		
610	615	620
Ile Lys Arg Lys Pro Pro Gly Cys Ser Asp Ser Pro Gly Val		
625	630	635
<210> 73		
<211> 639		
<212> PRT		
<213> Homo sapiens		
<400> 73		
Pro Glu Glu Gln Pro His Val Gly Asn Tyr Arg Leu Leu Arg Thr Ile		
1	5	10
15		
Gly Lys Gly Asn Ser Ala Lys Val Lys Leu Ala Arg His Ile Leu Thr		
20	25	30
Gly Arg Glu Val Ala Ile Lys Ile Ile Asp Lys Thr Gln Leu Asn Pro		
35	40	45
Ser Ser Leu Gln Lys Leu Phe Arg Glu Val Arg Ile Met Lys Gly Leu		
50	55	60

Asn	His	Pro	Asn	Ile	Val	Lys	Leu	Phe	Glu	Val	Ile	Glu	Thr	Glu	Lys
65					70						75				80
Thr	Leu	Tyr	Leu	Val	Met	Glu	Tyr	Ala	Ser	Ala	Gly	Glu	Val	Phe	Asp
					85				90					95	
Tyr	Leu	Val	Ser	His	Gly	Arg	Met	Lys	Glu	Lys	Glu	Ala	Arg	Ala	Lys
				100				105				110			
Phe	Arg	Gln	Ile	Val	Ser	Ala	Val	His	Tyr	Cys	His	Gln	Lys	Asn	Ile
					115			120				125			
Val	His	Arg	Asp	Leu	Lys	Ala	Glu	Asn	Leu	Leu	Asp	Ala	Glu	Ala	
					130			135			140				
Asn	Ile	Lys	Ile	Ala	Asp	Phe	Gly	Phe	Ser	Asn	Glu	Phe	Thr	Leu	Gly
					145			150			155			160	
Ser	Lys	Leu	Asp	Thr	Phe	Cys	Gly	Ser	Pro	Pro	Tyr	Ala	Ala	Pro	Glu
					165			170				175			
Leu	Phe	Gln	Gly	Lys	Lys	Tyr	Asp	Gly	Pro	Glu	Val	Asp	Ile	Trp	Ser
				180			185				190				
Leu	Gly	Val	Ile	Leu	Tyr	Thr	Leu	Val	Ser	Gly	Ser	Leu	Pro	Phe	Asp
					195			200			205				
Gly	His	Asn	Leu	Lys	Glu	Leu	Arg	Glu	Arg	Val	Leu	Arg	Gly	Lys	Tyr
					210			215			220				
Arg	Val	Pro	Phe	Tyr	Met	Ser	Thr	Asp	Cys	Glu	Ser	Ile	Leu	Arg	Arg
					225			230			235			240	
Phe	Leu	Val	Leu	Asn	Pro	Ala	Lys	Arg	Cys	Thr	Leu	Glu	Gln	Ile	Met
					245			250			255				
Lys	Asp	Lys	Trp	Ile	Asn	Ile	Gly	Tyr	Glu	Gly	Glu	Glu	Leu	Lys	Pro
					260			265			270				
Tyr	Thr	Glu	Pro	Glu	Glu	Asp	Phe	Gly	Asp	Thr	Lys	Arg	Ile	Glu	Val
					275			280			285				
Met	Val	Gly	Met	Gly	Tyr	Thr	Arg	Glu	Glu	Ile	Lys	Glu	Ser	Leu	Thr
					290			295			300				
Ser	Gln	Lys	Tyr	Asn	Glu	Val	Thr	Ala	Thr	Tyr	Leu	Leu	Gly	Arg	
					305			310			315			320	
Lys	Thr	Glu	Glu	Gly	Gly	Asp	Arg	Gly	Ala	Pro	Gly	Leu	Ala	Leu	Ala
					325			330			335				
Arg	Val	Arg	Ala	Pro	Ser	Asp	Thr	Thr	Asn	Gly	Thr	Ser	Ser	Ser	Lys
					340			345			350				
Gly	Thr	Ser	His	Ser	Lys	Gly	Gln	Arg	Ser	Ser	Ser	Ser	Thr	Tyr	His
					355			360			365				

Arg Gln Arg Arg His Ser Asp Phe Cys Gly Pro Ser Pro Ala Pro Leu
 370 375 380
 His Pro Lys Arg Ser Pro Thr Ser Thr Gly Glu Ala Glu Leu Lys Glu
 385 390 395 400
 Glu Arg Leu Pro Gly Arg Lys Ala Ser Cys Ser Thr Ala Gly Ser Gly
 405 410 415
 Ser Arg Gly Leu Pro Pro Ser Ser Pro Met Val Ser Ser Ala His Asn
 420 425 430
 Pro Asn Lys Ala Glu Ile Pro Glu Arg Arg Lys Asp Ser Thr Ser Thr
 435 440 445
 Pro Asn Asn Leu Pro Pro Ser Met Met Thr Arg Arg Asn Thr Tyr Val
 450 455 460
 Cys Thr Glu Arg Pro Gly Ala Glu Arg Pro Ser Leu Leu Pro Asn Gly
 465 470 475 480
 Lys Glu Asn Ser Ser Gly Thr Pro Arg Val Pro Pro Ala Ser Pro Ser
 485 490 495
 Ser His Ser Leu Ala Pro Pro Ser Gly Glu Arg Ser Arg Leu Ala Arg
 500 505 510
 Gly Ser Thr Ile Arg Ser Thr Phe His Gly Gly Gln Val Arg Asp Arg
 515 520 525
 Arg Ala Gly Gly Gly Gly Gly Val Gln Asn Gly Pro Pro Ala
 530 535 540
 Ser Pro Thr Leu Ala His Glu Ala Ala Pro Leu Pro Ala Gly Arg Pro
 545 550 555 560
 Arg Pro Thr Thr Asn Leu Phe Thr Lys Leu Thr Ser Lys Leu Thr Arg
 565 570 575
 Arg Val Thr Leu Asp Pro Ser Lys Arg Gln Asn Ser Asn Arg Cys Val
 580 585 590
 Ser Gly Ala Ser Leu Pro Gln Gly Ser Lys Ile Arg Ser Gln Thr Asn
 595 600 605
 Leu Arg Glu Ser Gly Asp Leu Arg Ser Gln Val Ala Ile Tyr Leu Gly
 610 615 620
 Ile Lys Arg Lys Pro Pro Gly Cys Ser Asp Ser Pro Gly Val
 625 630 635

<210> 74
 <211> 667
 <212> PRT
 <213> Homo sapiens

<400> 74
 Ala Asp Glu Gln Pro His Ile Gly Asn Tyr Arg Leu Leu Lys Thr Ile
 1 5 10 15
 Gly Lys Gly Asn Phe Ala Lys Val Lys Leu Ala Arg His Ile Leu Thr
 20 25 30
 Gly Arg Glu Val Ala Ile Lys Ile Ile Asp Lys Thr Gln Leu Asn Pro
 35 40 45
 Thr Ser Leu Gln Lys Leu Phe Arg Glu Val Arg Ile Met Lys Ile Leu
 50 55 60
 Asn His Pro Asn Ile Val Lys Leu Phe Glu Val Ile Glu Thr Gln Lys
 65 70 75 80
 Thr Leu Tyr Leu Ile Met Glu Tyr Ala Ser Gly Gly Lys Val Phe Asp
 85 90 95
 Tyr Leu Val Ala His Gly Arg Met Lys Glu Lys Glu Ala Arg Ser Lys
 100 105 110
 Phe Arg Gln Ile Val Ser Ala Val Gln Tyr Cys His Gln Lys Arg Ile
 115 120 125
 Val His Arg Asp Leu Lys Ala Glu Asn Leu Leu Asp Ala Asp Met
 130 135 140
 Asn Ile Lys Ile Ala Asp Phe Gly Phe Ser Asn Glu Phe Thr Val Gly
 145 150 155 160
 Gly Lys Leu Asp Thr Phe Cys Gly Ser Pro Pro Tyr Ala Ala Pro Glu
 165 170 175
 Leu Phe Gln Gly Lys Lys Tyr Asp Gly Pro Glu Val Asp Val Trp Ser
 180 185 190
 Leu Gly Val Ile Leu Tyr Thr Leu Val Ser Gly Ser Leu Pro Phe Asp
 195 200 205
 Gly Gln Asn Leu Lys Glu Leu Arg Glu Arg Val Leu Arg Gly Lys Tyr
 210 215 220
 Arg Ile Pro Phe Tyr Met Ser Thr Asp Cys Glu Asn Leu Leu Lys Arg
 225 230 235 240
 Phe Leu Val Leu Asn Pro Ile Lys Arg Gly Thr Leu Glu Gln Ile Met
 245 250 255
 Lys Asp Arg Trp Ile Asn Ala Gly His Glu Glu Asp Glu Leu Lys Pro
 260 265 270
 Phe Val Glu Pro Glu Leu Asp Ile Ser Asp Gln Lys Arg Ile Asp Ile
 275 280 285
 Met Val Gly Met Gly Tyr Ser Gln Glu Glu Ile Gln Glu Ser Leu Ser
 290 295 300

Lys Met Lys Tyr Asp Glu Ile Thr Ala Thr Tyr Leu Leu Leu Gly Arg
 305 310 315 320

 Lys Ser Ser Glu Val Arg Pro Ser Ser Asp Leu Asn Asn Ser Thr Gly
 325 330 335

 Gln Ser Pro His His Lys Val Gln Arg Ser Val Ser Ser Gln Lys
 340 345 350

 Gln Arg Arg Tyr Ser Asp His Ala Gly Pro Gly Ile Pro Ser Val Val
 355 360 365

 Ala Tyr Pro Lys Arg Ser Gln Thr Ser Thr Ala Asp Ser Asp Leu Lys
 370 375 380

 Glu Asp Gly Ile Ser Ser Arg Lys Ser Thr Gly Ser Ala Val Gly Gly
 385 390 395 400

 Lys Gly Ile Ala Pro Ala Ser Pro Met Leu Gly Asn Ala Ser Asn Pro
 405 410 415

 Asn Lys Ala Asp Ile Pro Glu Arg Lys Lys Ser Ser Thr Val Pro Ser
 420 425 430

 Ser Asn Thr Ala Ser Gly Gly Met Thr Arg Arg Asn Thr Tyr Val Cys
 435 440 445

 Ser Glu Arg Thr Thr Asp Asp Arg His Ser Val Ile Gln Asn Gly Lys
 450 455 460

 Glu Asn Ser Thr Ile Pro Asp Gln Arg Thr Pro Val Ala Ser Thr His
 465 470 475 480

 Ser Ile Ser Ser Ala Ala Thr Pro Asp Arg Ile Arg Phe Pro Arg Gly
 485 490 495

 Thr Ala Ser Arg Ser Thr Phe His Gly Gln Pro Arg Glu Arg Arg Thr
 500 505 510

 Ala Thr Tyr Asn Gly Pro Pro Ala Ser Pro Ser Leu Ser His Glu Ala
 515 520 525

 Thr Pro Leu Ser Gln Thr Arg Ser Arg Gly Ser Thr Thr Leu Phe Ser
 530 535 540

 Lys Leu Thr Ser Lys Leu Thr Arg Ser Arg Asn Val Ser Ala Lys Gln
 545 550 555 560

 Lys Asp Glu Asn Lys Glu Ala Lys Pro Arg Ser Leu Arg Phe Thr Trp
 565 570 575

 Ser Met Lys Thr Thr Ser Ser Met Asp Pro Gly Asp Met Met Arg Glu
 580 585 590

 Ile Arg Lys Val Leu Asp Ala Asn Asn Cys Asp Tyr Glu Gln Arg Glu
 595 600 605

Arg	Phe	Leu	Leu	Phe	Cys	Val	His	Gly	Asp	Gly	His	Ala	Glu	Asn	Leu
610						615						620			
Val	Gln	Trp	Glu	Met	Glu	Val	Cys	Lys	Leu	Pro	Arg	Leu	Ser	Leu	Asn
625					630				635			640			
Gly	Val	Arg	Phe	Lys	Arg	Ile	Ser	Gly	Thr	Ser	Ile	Ala	Phe	Lys	Asn
						645		650				655			
Ile	Ala	Ser	Lys	Ile	Ala	Asn	Glu	Leu	Lys	Leu					
						660		665							

<210> 75
<211> 888
<212> PRT
<213> Homo sapiens

<400> 75															
Met	Pro	Leu	Ala	Asn	His	Arg	Asp	Asp	Glu	His	Gly	Val	Ala	Ser	Met
1												10			15
Val	Ser	Val	His	Val	Glu	His	Pro	Gln	Glu	Ala	Ser	Val	Val	Val	His
												20		25	30
Gln	Val	Glu	Arg	Val	Ser	Gly	Pro	Trp	Glu	Glu	Ala	Asp	Ala	Glu	Ala
												35		40	45
Val	Ala	Arg	Ala	Glu	Ala	Ala	Ala	Arg	Ala	Glu	Ala	Ala	Ala	Pro	Tyr
												50		55	60
Thr	Val	Leu	Ala	Gln	Ser	Ala	Pro	Arg	Glu	Asp	Gly	Tyr	Ser	Asp	Ala
												65		70	75
Ser	Gly	Phe	Gly	Tyr	Cys	Phe	Arg	Glu	Leu	Arg	Gly	Gly	Glu	Cys	Ala
												85		90	95
Ser	Pro	Leu	Pro	Gly	Leu	Arg	Thr	Gln	Glu	Val	Cys	Cys	Arg	Gly	Ala
												100		105	110
Gly	Leu	Ala	Trp	Gly	Val	His	Asp	Cys	Gln	Leu	Cys	Ser	Glu	Arg	Leu
												115		120	125
Gly	Asn	Ser	Glu	Arg	Val	Ser	Ala	Pro	Asp	Gly	Pro	Cys	Pro	Thr	Gly
												130		135	140
Phe	Glu	Arg	Val	Asn	Gly	Ser	Cys	Glu	Asp	Val	Asp	Glu	Cys	Ala	Thr
												145		150	155
Gly	Gly	Arg	Cys	Gln	His	Gly	Glu	Cys	Ala	Asn	Thr	Arg	Gly	Gly	Tyr
												165		170	175
Thr	Cys	Val	Cys	Pro	Asp	Gly	Phe	Leu	Leu	Asp	Ser	Ser	Arg	Ser	Ser
												180		185	190
Cys	Ile	Ser	Gln	His	Val	Ile	Ser	Glu	Ala	Lys	Gly	Pro	Cys	Phe	Arg

195	200	205
Val Leu Arg Asp Gly Gly Cys Ser	Leu Pro Ile Leu Arg Asn Ile Thr	
210	215	220
Lys Gln Ile Cys Cys Cys Ser Arg Val Gly Lys Ala Trp Gly Arg Gly		
225	230	235 240
Cys Gln Leu Cys Pro Pro Phe Gly Ser Glu Gly Phe Arg Glu Ile Cys		
245	250	255
Pro Ala Gly Pro Gly Tyr His Tyr Ser Ala Ser Asp Leu Arg Tyr Asn		
260	265	270
Thr Arg Pro Leu Gly Gln Glu Pro Pro Arg Val Ser Leu Ser Gln Pro		
275	280	285
Arg Thr Leu Pro Ala Thr Ser Arg Pro Ser Ala Gly Phe Leu Pro Thr		
290	295	300
His Arg Leu Glu Pro Arg Pro Glu Pro Arg Pro Asp Pro Arg Pro Gly		
305	310	315 320
Pro Glu Leu Pro Leu Pro Ser Ile Pro Ala Trp Thr Gly Pro Glu Ile		
325	330	335
Pro Glu Ser Gly Pro Ser Ser Gly Met Cys Gln Arg Asn Pro Gln Val		
340	345	350
Cys Gly Pro Gly Arg Cys Ile Ser Arg Pro Ser Gly Tyr Thr Cys Ala		
355	360	365
Cys Asp Ser Gly Phe Arg Leu Ser Pro Gln Gly Thr Arg Cys Ile Asp		
370	375	380
Val Asp Glu Cys Arg Arg Val Pro Pro Cys Ala Pro Gly Arg Cys		
385	390	395 400
Glu Asn Ser Pro Gly Ser Phe Arg Cys Val Cys Gly Pro Gly Phe Arg		
405	410	415
Ala Gly Pro Arg Ala Ala Glu Cys Leu Asp Val Asp Glu Cys His Arg		
420	425	430
Val Pro Pro Pro Cys Asp Leu Gly Arg Cys Glu Asn Thr Pro Gly Ser		
435	440	445
Phe Leu Cys Val Cys Pro Ala Gly Tyr Gln Ala Ala Pro His Gly Ala		
450	455	460
Ser Cys Gln Asp Val Asp Glu Cys Thr Gln Ser Pro Gly Leu Cys Gly		
465	470	475 480
Arg Gly Ala Cys Lys Asn Leu Pro Gly Ser Phe Arg Cys Val Cys Pro		
485	490	495
Ala Gly Phe Arg Gly Ser Ala Cys Glu Glu Asp Val Asp Glu Cys Ala		

500	505	510
Gln Glu Pro Pro Pro Cys Gly Pro Gly Arg Cys Asp Asn Thr Ala Gly		
515	520	525
Ser Phe His Cys Ala Cys Pro Ala Gly Phe Arg Ser Arg Gly Pro Gly		
530	535	540
Ala Pro Cys Gln Asp Val Asp Glu Cys Ala Arg Ser Pro Pro Pro Cys		
545	550	555
Thr Tyr Gly Arg Cys Glu Asn Thr Glu Gly Ser Phe Gln Cys Val Cys		
565	570	575
Pro Met Gly Phe Gln Pro Asn Thr Ala Gly Ser Glu Cys Glu Asp Val		
580	585	590
Asp Glu Cys Glu Asn His Leu Ala Cys Pro Gly Gln Glu Cys Val Asn		
595	600	605
Ser Pro Gly Ser Phe Gln Cys Arg Thr Cys Pro Ser Gly His His Leu		
610	615	620
His Arg Gly Arg Cys Thr Asp Val Asp Glu Cys Ser Ser Gly Ala Pro		
625	630	635
Pro Cys Gly Pro His Gly His Cys Thr Asn Thr Glu Gly Ser Phe Arg		
645	650	655
Cys Ser Cys Ala Pro Gly Tyr Arg Ala Pro Ser Gly Arg Pro Gly Pro		
660	665	670
Cys Ala Asp Val Asn Glu Cys Leu Glu Gly Asp Phe Cys Phe Pro His		
675	680	685
Gly Glu Cys Leu Asn Thr Asp Gly Ser Phe Ala Cys Thr Cys Ala Pro		
690	695	700
Gly Tyr Arg Pro Gly Pro Arg Gly Ala Ser Cys Leu Asp Val Asp Glu		
705	710	715
Cys Ser Glu Glu Asp Leu Cys Gln Ser Gly Ile Cys Thr Asn Thr Asp		
725	730	735
Gly Ser Phe Glu Cys Ile Cys Pro Pro Gly His Arg Ala Gly Pro Asp		
740	745	750
Leu Ala Ser Cys Leu Asp Val Asp Glu Cys Arg Glu Arg Gly Pro Ala		
755	760	765
Leu Cys Gly Ser Gln Arg Cys Glu Asn Ser Pro Gly Ser Tyr Arg Cys		
770	775	780
Val Arg Asp Cys Asp Pro Gly Tyr His Ala Gly Pro Glu Gly Thr Cys		
785	790	795
Asp Asp Val Asp Glu Cys Gln Glu Tyr Gly Pro Glu Ile Cys Gly Ala		

805	810	815
Gln Arg Cys Glu Asn Thr Pro Gly Ser Tyr Arg Cys Thr Pro Ala Cys		
820	825	830
Asp Pro Gly Tyr Gln Pro Thr Pro Gly Gly Cys Gln Asp Val Asp		
835	840	845
Glu Cys Arg Asn Arg Ser Phe Cys Gly Ala His Ala Val Cys Gln Asn		
850	855	860
Leu Pro Gly Ser Phe Gln Cys Leu Cys Asp Gln Val Thr Arg Gly His		
865	870	875
Gly Met Gly Val Thr Ala Trp Met		
885		
<210> 76		
<211> 1511		
<212> PRT		
<213> Homo sapiens		
<400> 76		
Met Gly Arg Pro Ala Pro Ala Val Pro Arg Pro Ala Arg Pro Ala Thr		
1	5	10
15		
Pro Pro Ala Trp Thr Ala Ala Leu Pro Ala Gly Arg Pro Arg Gly Asp		
20	25	30
Pro Gly Phe Arg Ala Phe Leu Cys Pro Leu Ile Cys His Asn Gly Gly		
35	40	45
Val Cys Val Lys Pro Asp Arg Cys Leu Cys Pro Pro Asp Phe Ala Gly		
50	55	60
Lys Phe Cys Gln Leu His Ser Ser Gly Ala Arg Pro Pro Ala Pro Ala		
65	70	75
80		
Ile Pro Gly Leu Thr Arg Ser Val Tyr Thr Met Pro Leu Ala Asn His		
85	90	95
Arg Asp Asp Glu His Gly Val Ala Ser Met Val Ser Val His Val Glu		
100	105	110
His Pro Gln Glu Ala Ser Val Val Val His Gln Val Glu Arg Val Ser		
115	120	125
Gly Pro Trp Glu Glu Ala Asp Ala Glu Ala Val Ala Arg Ala Glu Ala		
130	135	140
Ala Ala Arg Ala Glu Ala Ala Ala Pro Tyr Thr Val Leu Ala Gln Ser		
145	150	155
160		
Ala Pro Arg Glu Asp Gly Tyr Ser Asp Ala Ser Gly Phe Gly Tyr Cys		
165	170	175

Phe Arg Glu Leu Arg Gly Gly Glu Cys Ala Ser Pro Leu Pro Gly Leu
180 185 190

Arg Thr Gln Glu Val Cys Cys Arg Gly Ala Gly Leu Ala Trp Gly Val
195 200 205

His Asp Cys Gln Leu Cys Ser Glu Arg Leu Gly Asn Ser Glu Arg Val
210 215 220

Ser Ala Pro Asp Gly Pro Cys Pro Thr Gly Phe Glu Arg Val Asn Gly
225 230 235 240

Ser Cys Glu Asp Val Asp Glu Cys Ala Thr Gly Gly Arg Cys Gln His
245 250 255

Gly Glu Cys Ala Asn Thr Arg Gly Gly Tyr Thr Cys Val Cys Pro Asp
260 265 270

Gly Phe Leu Leu Asp Ser Ser Arg Ser Ser Cys Ile Ser Gln His Val
275 280 285

Ile Ser Glu Ala Lys Gly Pro Cys Phe Arg Val Leu Arg Asp Gly Gly
290 295 300

Cys Ser Leu Pro Ile Leu Arg Asn Ile Thr Lys Gln Ile Cys Cys Cys
305 310 315 320

Ser Arg Val Gly Lys Ala Trp Gly Arg Gly Cys Gln Leu Cys Pro Pro
325 330 335

Phe Gly Ser Glu Gly Phe Arg Glu Ile Cys Pro Ala Gly Pro Gly Tyr
340 345 350

His Tyr Ser Ala Ser Asp Leu Arg Tyr Asn Thr Arg Pro Leu Gly Gln
355 360 365

Glu Pro Pro Arg Val Ser Leu Ser Gln Pro Arg Thr Leu Pro Ala Thr
370 375 380

Ser Arg Pro Ser Ala Gly Phe Leu Pro Thr His Arg Leu Glu Pro Arg
385 390 395 400

Pro Glu Pro Arg Pro Asp Pro Arg Pro Gly Pro Glu Phe Pro Leu Pro
405 410 415

Ser Ile Pro Ala Trp Thr Gly Pro Glu Ile Pro Glu Ser Gly Pro Ser
420 425 430

Ser Gly Met Cys Gln Arg Asn Pro Gln Val Cys Gly Pro Gly Arg Cys
435 440 445

Ile Ser Arg Pro Ser Gly Tyr Thr Cys Ala Cys Asp Ser Gly Phe Arg
450 455 460

Leu Ser Pro Gln Gly Thr Arg Cys Ile Asp Val Asp Glu Cys Arg Arg
465 470 475 480

Val Pro Pro Pro Cys Ala Pro Gly Arg Cys Glu Asn Ser Pro Gly Ser
 485 490 495

 Phe Arg Cys Val Cys Gly Pro Gly Phe Arg Ala Gly Pro Arg Ala Ala
 500 505 510

 Glu Cys Leu Asp Val Asp Glu Cys His Arg Val Pro Pro Pro Cys Asp
 515 520 525

 Leu Gly Arg Cys Glu Asn Thr Pro Gly Ser Phe Leu Cys Val Cys Pro
 530 535 540

 Ala Gly Tyr Gln Ala Ala Pro His Gly Ala Ser Cys Gln Asp Val Asp
 545 550 555 560

 Glu Cys Thr Gln Ser Pro Gly Leu Cys Gly Arg Gly Ala Cys Lys Asn
 565 570 575

 Leu Pro Gly Ser Phe Arg Cys Val Cys Pro Ala Gly Phe Arg Gly Ser
 580 585 590

 Ala Cys Glu Glu Asp Val Asp Glu Cys Ala Gln Glu Pro Pro Pro Cys
 595 600 605

 Gly Pro Gly Arg Cys Asp Asn Thr Ala Gly Ser Phe His Cys Ala Cys
 610 615 620

 Pro Ala Gly Phe Arg Ser Arg Gly Pro Gly Ala Pro Cys Gln Asp Val
 625 630 635 640

 Asp Glu Cys Ala Arg Ser Pro Pro Pro Cys Thr Tyr Gly Arg Cys Glu
 645 650 655

 Asn Thr Glu Gly Ser Phe Gln Cys Val Cys Pro Met Gly Phe Gln Pro
 660 665 670

 Asn Ala Ala Gly Ser Glu Cys Glu Asp Val Asp Glu Cys Glu Asn His
 675 680 685

 Leu Ala Cys Pro Gly Gln Glu Cys Val Asn Ser Pro Gly Ser Phe Gln
 690 695 700

 Cys Arg Ala Cys Pro Ser Gly His His Leu His Arg Gly Arg Cys Thr
 705 710 715 720

 Asp Val Asp Glu Cys Ser Ser Gly Ala Pro Pro Cys Gly Pro His Gly
 725 730 735

 His Cys Thr Asn Thr Glu Gly Ser Phe Arg Cys Ser Cys Ala Pro Gly
 740 745 750

 Tyr Arg Ala Pro Ser Gly Arg Pro Gly Pro Cys Ala Asp Val Asn Glu
 755 760 765

 Cys Leu Glu Gly Asp Phe Cys Phe Pro His Gly Glu Cys Leu Asn Thr
 770 775 780

Asp Gly Ser Phe Ala Cys Thr Cys Ala Pro Gly Tyr Arg Pro Gly Pro
 785 790 795 800
 Arg Gly Ala Ser Cys Leu Asp Val Asp Glu Cys Ser Glu Glu Asp Leu
 805 810 815
 Cys Gln Ser Gly Ile Cys Thr Asn Thr Asp Gly Ser Phe Glu Cys Ile
 820 825 830
 Cys Pro Pro Gly His Arg Ala Gly Pro Asp Leu Ala Ser Cys Leu Asp
 835 840 845
 Val Asp Glu Cys Arg Glu Arg Gly Pro Ala Leu Cys Gly Ser Gln Arg
 850 855 860
 Cys Glu Asn Ser Pro Gly Ser Tyr Arg Cys Val Arg Asp Cys Asp Pro
 865 870 875 880
 Gly Tyr His Ala Gly Pro Glu Gly Thr Cys Asp Asp Val Asp Glu Cys
 885 890 895
 Gln Glu Tyr Gly Pro Glu Ile Cys Gly Ala Gln Arg Cys Glu Asn Thr
 900 905 910
 Pro Gly Ser Tyr Arg Cys Thr Pro Ala Cys Asp Pro Gly Tyr Gln Pro
 915 920 925
 Thr Pro Gly Gly Cys Gln Asp Val Asp Glu Cys Arg Asn Arg Ser
 930 935 940
 Phe Cys Gly Ala His Ala Val Cys Gln Asn Leu Pro Gly Ser Phe Gln
 945 950 955 960
 Cys Leu Cys Asp Gln Gly Tyr Glu Gly Ala Arg Asp Gly Arg His Cys
 965 970 975
 Val Asp Val Asn Glu Cys Glu Thr Leu Gln Gly Val Cys Gly Ala Ala
 980 985 990
 Leu Cys Glu Asn Val Glu Gly Ser Phe Leu Cys Val Cys Pro Asn Ser
 995 1000 1005
 Pro Glu Glu Phe Asp Pro Met Thr Gly Arg Cys Val Pro Pro Arg Thr
 1010 1015 1020
 Ser Ala Gly Met Phe Pro Gly Ser Gln Pro Gln Ala Pro Ala Ser Pro
 1025 1030 1035 1040
 Val Leu Pro Ala Arg Pro Pro Pro Pro Leu Pro Arg Arg Pro Ser
 1045 1050 1055
 Thr Pro Arg Gln Gly Pro Val Gly Ser Gly Arg Arg Glu Cys Tyr Phe
 1060 1065 1070
 Asp Thr Ala Ala Pro Asp Ala Cys Asp Asn Ile Leu Ala Arg Asn Val
 1075 1080 1085

Thr Trp Gln Glu Cys Cys Cys Thr Val Gly Glu Gly Trp Gly Ser Gly
1090 1095 1100

Cys Arg Ile Gln Gln Cys Pro Gly Thr Glu Thr Ala Glu Tyr Gln Ser
1105 1110 1115 1120

Leu Cys Pro His Gly Arg Gly Tyr Leu Ala Pro Ser Gly Asp Leu Ser
1125 1130 1135

Leu Arg Arg Asp Val Asp Glu Cys Gln Leu Phe Arg Asp Gln Val Cys
1140 1145 1150

Lys Ser Gly Val Cys Val Asn Thr Ala Pro Gly Tyr Ser Cys Tyr Cys
1155 1160 1165

Ser Asn Gly Tyr Tyr Tyr His Thr Gln Arg Leu Glu Cys Ile Asp Asn
1170 1175 1180

Asp Glu Cys Ala Asp Glu Glu Pro Ala Cys Glu Gly Gly Arg Cys Val
1185 1190 1195 1200

Asn Thr Val Gly Ser Tyr His Cys Thr Cys Glu Pro Pro Leu Val Leu
1205 1210 1215

Asp Gly Ser Gln Arg Arg Cys Val Ser Asn Glu Ser Gln Ser Leu Asp
1220 1225 1230

Asp Asn Leu Gly Val Cys Trp Gln Glu Val Gly Ala Asp Leu Val Cys
1235 1240 1245

Ser His Pro Arg Leu Asp Arg Gln Ala Thr Tyr Thr Glu Cys Cys Cys
1250 1255 1260

Leu Tyr Gly Glu Ala Trp Gly Met Asp Cys Ala Leu Cys Pro Ala Gln
1265 1270 1275 1280

Asp Ser Asp Asp Phe Glu Ala Leu Cys Asn Val Leu Arg Pro Pro Ala
1285 1290 1295

Tyr Ser Pro Pro Arg Pro Gly Gly Phe Gly Leu Pro Tyr Glu Tyr Gly
1300 1305 1310

Pro Asp Leu Gly Pro Pro Tyr Gln Gly Leu Pro Tyr Gly Pro Glu Leu
1315 1320 1325

Tyr Pro Pro Pro Ala Leu Pro Tyr Asp Pro Tyr Pro Pro Pro Pro Gly
1330 1335 1340

Pro Phe Ala Arg Arg Glu Ala Pro Tyr Gly Ala Pro Arg Phe Asp Met
1345 1350 1355 1360

Pro Asp Phe Glu Asp Asp Gly Gly Pro Tyr Gly Glu Ser Glu Ala Pro
1365 1370 1375

Ala Pro Pro Gly Pro Gly Thr Arg Trp Pro Tyr Arg Ser Arg Asp Thr
1380 1385 1390

Arg Arg Ser Phe Pro Glu Pro Glu Glu Pro Pro Glu Gly Gly Ser Tyr
1395 1400 1405

Ala Gly Ser Leu Ala Glu Pro Tyr Glu Glu Leu Glu Ala Glu Glu Cys
1410 1415 1420

Gly Ile Leu Asp Gly Cys Thr Asn Gly Arg Cys Val Arg Val Pro Glu
1425 1430 1435 1440

Gly Phe Thr Cys Arg Cys Phe Asp Gly Tyr Arg Leu Asp Met Thr Arg
1445 1450 1455

Met Ala Cys Val Asp Ile Asn Glu Cys Asp Glu Ala Glu Ala Ala Ser
1460 1465 1470

Pro Leu Cys Val Asn Ala Arg Cys Leu Asn Thr Asp Gly Ser Phe Arg
1475 1480 1485

Cys Ile Cys Arg Pro Gly Phe Ala Pro Thr His Gln Pro His His Cys
1490 1495 1500

Ala Pro Ala Arg Pro Arg Ala
1505 1510

<210> 77
<211> 1587
<212> PRT
<213> Homo sapiens

<400> 77
Met Gly Asp Val Lys Ala Leu Leu Phe Val Val Ala Ala Arg Ala Arg
1 5 10 15

Arg Leu Gly Gly Ala Ala Ala Ser Glu Ser Leu Ala Val Ser Glu Ala
20 25 30

Phe Cys Arg Val Arg Ser Cys Gln Pro Lys Lys Cys Ala Gly Pro Gln
35 40 45

Arg Cys Leu Asn Pro Val Pro Ala Val Pro Ser Pro Ser Pro Ser Val
50 55 60

Arg Lys Arg Gln Val Ser Leu Asn Trp Gln Pro Leu Thr Leu Gln Glu
65 70 75 80

Ala Arg Ala Leu Leu Lys Arg Arg Arg Pro Arg Gly Pro Gly Gly Arg
85 90 95

Gly Leu Leu Arg Arg Arg Pro Pro Gln Arg Ala Pro Ala Gly Lys Ala
100 105 110

Pro Val Leu Cys Pro Leu Ile Cys His Asn Gly Gly Val Cys Val Lys
115 120 125

Pro Asp Arg Cys Phe Cys Pro Pro Asp Phe Ala Gly Lys Phe Cys Gln
130 135 140

Leu His Ser Ser Gly Ala Arg Pro Pro Ala Pro Ala Val Pro Gly Leu
 145 150 155 160
 Thr Arg Ser Val Tyr Thr Met Pro Leu Ala Asn His Arg Asp Asp Glu
 165 170 175
 His Gly Val Ala Ser Met Val Ser Val His Val Glu His Pro Gln Glu
 180 185 190
 Ala Ser Val Val Val His Gln Val Glu Arg Val Ser Gly Pro Trp Glu
 195 200 205
 Glu Ala Asp Ala Glu Ala Val Ala Arg Ala Glu Ala Ala Ala Arg Ala
 210 215 220
 Glu Ala Ala Ala Pro Tyr Thr Val Leu Ala Gln Ser Ala Pro Arg Glu
 225 230 235 240
 Asp Gly Tyr Ser Asp Ala Ser Gly Phe Gly Tyr Cys Phe Arg Glu Leu
 245 250 255
 Arg Gly Gly Glu Cys Ala Ser Pro Leu Pro Gly Leu Arg Thr Gln Glu
 260 265 270
 Val Cys Cys Arg Gly Ala Gly Leu Ala Trp Gly Val His Asp Cys Gln
 275 280 285
 Leu Cys Ser Glu Arg Leu Gly Asn Ser Glu Arg Val Ser Ala Pro Asp
 290 295 300
 Gly Pro Cys Pro Thr Gly Phe Glu Arg Val Asn Gly Ser Cys Glu Asp
 305 310 315 320
 Val Asp Glu Cys Ala Thr Gly Gly Arg Cys Gln His Gly Glu Cys Ala
 325 330 335
 Asn Thr Arg Gly Gly Tyr Thr Cys Val Cys Pro Asp Gly Phe Leu Leu
 340 345 350
 Asp Ser Ser Arg Ser Ser Cys Ile Ser Gln His Val Ile Ser Glu Ala
 355 360 365
 Lys Gly Pro Cys Phe Arg Val Leu Arg Asp Gly Gly Cys Ser Leu Pro
 370 375 380
 Ile Leu Arg Asn Ile Thr Lys Gln Ile Cys Cys Cys Ser Arg Val Gly
 385 390 395 400
 Lys Ala Trp Gly Arg Gly Cys Gln Leu Cys Pro Pro Phe Gly Ser Glu
 405 410 415
 Gly Phe Arg Glu Ile Cys Pro Ala Gly Pro Gly Tyr His Tyr Ser Ala
 420 425 430
 Ser Asp Leu Arg Tyr Asn Thr Arg Pro Leu Gly Gln Glu Pro Pro Arg
 435 440 445

Val Ser Leu Ser Gln Pro Arg Thr Leu Pro Ala Thr Ser Arg Pro Ser
 450 455 460
 Ala Gly Phe Leu Pro Thr His Arg Leu Glu Pro Arg Pro Glu Pro Arg
 465 470 475 480
 Pro Asp Pro Arg Pro Gly Pro Glu Leu Pro Leu Pro Ser Ile Pro Ala
 485 490 495
 Trp Thr Gly Pro Glu Ile Pro Glu Ser Gly Pro Ser Ser Gly Met Cys
 500 505 510
 Gln Arg Asn Pro Gln Val Cys Gly Pro Gly Arg Cys Ile Ser Arg Pro
 515 520 525
 Ser Gly Tyr Thr Cys Ala Cys Asp Ser Gly Phe Arg Leu Ser Pro Gln
 530 535 540
 Gly Thr Arg Cys Ile Asp Val Asp Glu Cys Arg Arg Val Pro Pro Pro
 545 550 555 560
 Cys Ala Pro Gly Arg Cys Glu Asn Ser Pro Gly Ser Phe Arg Cys Val
 565 570 575
 Cys Gly Pro Gly Phe Arg Ala Gly Pro Arg Ala Ala Glu Cys Leu Asp
 580 585 590
 Val Asp Glu Cys His Arg Val Pro Pro Pro Cys Asp Leu Gly Arg Cys
 595 600 605
 Glu Asn Thr Pro Gly Ser Phe Leu Cys Val Cys Pro Ala Gly Tyr Gln
 610 615 620
 Ala Ala Pro His Gly Ala Ser Cys Gln Asp Val Asp Glu Cys Thr Gln
 625 630 635 640
 Ser Pro Gly Leu Cys Gly Arg Gly Cys Lys Asn Leu Pro Gly Ser
 645 650 655
 Phe Arg Cys Val Cys Pro Ala Gly Phe Arg Gly Ser Ala Cys Glu Glu
 660 665 670
 Asp Val Asp Glu Cys Ala Gln Glu Pro Pro Pro Cys Gly Pro Gly Arg
 675 680 685
 Cys Asp Asn Thr Ala Gly Ser Phe His Cys Ala Cys Pro Ala Gly Phe
 690 695 700
 Arg Ser Arg Gly Pro Gly Ala Pro Cys Gln Asp Val Asp Glu Cys Ala
 705 710 715 720
 Arg Ser Pro Pro Pro Cys Thr Tyr Gly Arg Cys Glu Asn Thr Glu Gly
 725 730 735
 Ser Phe Gln Cys Val Cys Pro Met Gly Phe Gln Pro Asn Thr Ala Gly
 740 745 750

Ser Glu Cys Glu Asp Val Asp Glu Cys Glu Asn His Leu Ala Cys Pro
 755 760 765
 Gly Gln Glu Cys Val Asn Ser Pro Gly Ser Phe Gln Cys Arg Thr Cys
 770 775 780
 Pro Ser Gly His His Leu His Arg Gly Arg Cys Thr Asp Val Asp Glu
 785 790 795 800
 Cys Ser Ser Gly Ala Pro Pro Cys Gly Pro His Gly His Cys Thr Asn
 805 810 815
 Thr Glu Gly Ser Phe Arg Cys Ser Cys Ala Pro Gly Tyr Arg Ala Pro
 820 825 830
 Ser Gly Arg Pro Gly Pro Cys Ala Asp Val Asn Glu Cys Leu Glu Gly
 835 840 845
 Asp Phe Cys Phe Pro His Gly Glu Cys Leu Asn Thr Asp Gly Ser Phe
 850 855 860
 Ala Cys Thr Cys Ala Pro Gly Tyr Arg Pro Gly Pro Arg Gly Ala Ser
 865 870 875 880
 Cys Leu Asp Val Asp Glu Cys Ser Glu Glu Asp Leu Cys Gln Ser Gly
 885 890 895
 Ile Cys Thr Asn Thr Asp Gly Ser Phe Glu Cys Ile Cys Pro Pro Gly
 900 905 910
 His Arg Ala Gly Pro Asp Leu Ala Ser Cys Leu Asp Val Asp Glu Cys
 915 920 925
 Arg Glu Arg Gly Pro Ala Leu Cys Gly Ser Gln Arg Cys Glu Asn Ser
 930 935 940
 Pro Gly Ser Tyr Arg Cys Val Arg Asp Cys Asp Pro Gly Tyr His Ala
 945 950 955 960
 Gly Pro Glu Gly Thr Cys Asp Asp Val Asp Glu Cys Gln Glu Tyr Gly
 965 970 975
 Pro Glu Ile Cys Gly Ala Gln Arg Cys Glu Asn Thr Pro Gly Ser Tyr
 980 985 990
 Arg Cys Thr Pro Ala Cys Asp Pro Gly Tyr Gln Pro Thr Pro Gly Gly
 995 1000 1005
 Gly Cys Gln Asp Val Asp Glu Cys Arg Asn Arg Ser Phe Cys Gly Ala
 1010 1015 1020
 His Ala Val Cys Gln Asn Leu Pro Gly Ser Phe Gln Cys Leu Cys Asp
 1025 1030 1035 1040
 Gln Gly Tyr Glu Gly Ala Arg Asp Gly Arg His Cys Val Asp Val Asn
 1045 1050 1055

Glu Cys Glu Thr Leu Gln Gly Val Cys Gly Ala Ala Leu Cys Glu Asn
 1060 1065 1070
 Val Glu Gly Ser Phe Leu Cys Val Cys Pro Asn Ser Pro Glu Glu Phe
 1075 1080 1085
 Asp Pro Met Thr Gly Arg Cys Val Pro Pro Arg Thr Ser Val Gly Met
 1090 1095 1100
 Ser Pro Gly Ser Gln Pro Gln Ala Pro Val Ser Pro Val Leu Pro Ala
 1105 1110 1115 1120
 Arg Pro Pro Pro Pro Leu Ser Arg Arg Pro Arg Lys Pro Arg Lys
 1125 1130 1135
 Gly Pro Val Gly Ser Gly Cys Arg Glu Cys Tyr Phe Asp Thr Ala Ala
 1140 1145 1150
 Pro Asp Ala Cys Asp Asn Ile Leu Ala Arg Asn Val Thr Trp Gln Glu
 1155 1160 1165
 Cys Cys Cys Thr Val Gly Glu Gly Trp Gly Ser Gly Cys Arg Ile Gln
 1170 1175 1180
 Gln Cys Pro Gly Thr Glu Thr Ala Glu Tyr Gln Ser Leu Cys Pro His
 1185 1190 1195 1200
 Gly Arg Gly Tyr Leu Ala Pro Ser Gly Asp Leu Ser Leu Arg Arg Asp
 1205 1210 1215
 Val Asp Glu Cys Gln Leu Phe Arg Asp Gln Val Cys Lys Ser Gly Val
 1220 1225 1230
 Cys Val Asn Thr Ala Pro Gly Tyr Ser Cys Tyr Cys Ser Asn Gly Tyr
 1235 1240 1245
 Tyr Tyr His Thr Gln Arg Leu Glu Cys Ile Asp Asn Asp Glu Cys Ala
 1250 1255 1260
 Asp Glu Glu Pro Ala Cys Glu Gly Gly Arg Cys Val Asn Thr Val Gly
 1265 1270 1275 1280
 Ser Tyr His Cys Thr Cys Glu Pro Pro Leu Val Leu Asp Gly Ser Gln
 1285 1290 1295
 Arg Arg Cys Val Ser Asn Glu Ser Gln Ser Leu Asp Asp Asn Leu Gly
 1300 1305 1310
 Val Cys Trp Gln Glu Val Gly Ala Asp Leu Val Cys Ser His Pro Arg
 1315 1320 1325
 Leu Asp Arg Gln Ala Thr Tyr Thr Glu Cys Cys Cys Leu Tyr Gly Glu
 1330 1335 1340
 Ala Trp Gly Met Asp Cys Ala Leu Cys Pro Ala Gln Asp Ser Asp Asp
 1345 1350 1355 1360

Phe Glu Ala Leu Cys Asn Val Leu Arg Pro Pro Ala Tyr Ser Pro Pro
 1365 1370 1375

 Arg Pro Gly Gly Phe Gly Leu Pro Tyr Glu Tyr Gly Pro Asp Leu Gly
 1380 1385 1390

 Pro Pro Tyr Gln Gly Leu Pro Tyr Gly Pro Glu Leu Tyr Pro Pro Pro
 1395 1400 1405

 Ala Leu Pro Tyr Asp Pro Tyr Pro Pro Pro Gly Pro Phe Ala Arg
 1410 1415 1420

 Arg Glu Ala Pro Tyr Gly Ala Pro Arg Phe Asp Met Pro Asp Phe Glu
 1425 1430 1440

 Asp Asp Gly Gly Pro Tyr Gly Glu Ser Glu Ala Pro Ala Pro Pro Gly
 1445 1450 1455

 Pro Gly Thr Arg Trp Pro Tyr Arg Ser Arg Asp Thr Arg Arg Ser Phe
 1460 1465 1470

 Pro Glu Pro Glu Glu Pro Pro Glu Gly Gly Ser Tyr Ala Gly Ser Leu
 1475 1480 1485

 Ala Glu Pro Tyr Glu Glu Leu Glu Ala Glu Glu Cys Gly Ile Leu Asp
 1490 1495 1500

 Gly Cys Thr Asn Asp Arg Cys Val Arg Val Pro Glu Gly Phe Thr Cys
 1505 1510 1520

 Arg Cys Phe Asp Gly Tyr Arg Leu Asp Met Thr Arg Met Ala Cys Val
 1525 1530 1535

 Asp Ile Asn Glu Cys Asp Glu Ala Glu Ala Ala Ser Pro Leu Cys Val
 1540 1545 1550

 Asn Ala Arg Cys Leu Asn Thr Asp Gly Ser Phe Arg Cys Ile Cys Arg
 1555 1560 1565

 Pro Gly Phe Ala Pro Thr His Gln Pro His His Cys Ala Pro Ala Arg
 1570 1575 1580

Pro Arg Ala
 1585

<210> 78
 <211> 775
 <212> PRT
 <213> Homo sapiens

<400> 78
 Met Pro Leu Ala Asn His Arg Asp Asp Glu His Gly Val Ala Ser Met
 1 5 10 15

 Val Ser Val His Val Glu His Pro Gln Glu Ala Ser Val Val Val His

20	25	30
Gln Val Glu Arg Val Ser Gly Pro Trp Glu Glu Ala Asp Ala Glu Ala		
35	40	45
Val Ala Arg Ala Glu Ala Ala Arg Ala Glu Ala Ala Ala Pro Tyr		
50	55	60
Thr Val Leu Ala Gln Ser Ala Pro Arg Glu Asp Gly Tyr Ser Asp Ala		
65	70	75
Ser Gly Phe Gly Tyr Cys Phe Arg Glu Leu Arg Gly Gly Glu Cys Ala		
85	90	95
Ser Pro Leu Pro Gly Leu Arg Thr Gln Glu Val Cys Cys Arg Gly Ala		
100	105	110
Gly Leu Ala Trp Gly Val His Asp Cys Gln Leu Cys Ser Glu Arg Leu		
115	120	125
Gly Asn Ser Glu Arg Val Ser Ala Pro Asp Gly Pro Cys Pro Thr Gly		
130	135	140
Phe Glu Arg Val Asn Gly Ser Cys Glu Asp Val Asp Glu Cys Ala Thr		
145	150	155
160		
Gly Gly Arg Cys Gln His Gly Glu Cys Ala Asn Thr Arg Gly Gly Tyr		
165	170	175
Thr Cys Val Cys Pro Asp Gly Phe Leu Leu Asp Ser Ser Arg Ser Ser		
180	185	190
Cys Ile Ser Gln His Val Ile Ser Glu Ala Lys Gly Pro Cys Phe Arg		
195	200	205
Val Leu Arg Asp Gly Gly Cys Ser Leu Pro Ile Leu Arg Asn Ile Thr		
210	215	220
Lys Gln Ile Cys Cys Cys Ser Arg Val Gly Lys Ala Trp Gly Arg Gly		
225	230	235
240		
Cys Gln Leu Cys Pro Pro Phe Gly Ser Glu Gly Phe Arg Glu Ile Cys		
245	250	255
Pro Ala Gly Pro Gly Tyr His Tyr Ser Ala Ser Asp Leu Arg Tyr Asn		
260	265	270
Thr Arg Pro Leu Gly Gln Glu Pro Pro Arg Val Ser Leu Ser Gln Pro		
275	280	285
Arg Thr Leu Pro Ala Thr Ser Arg Pro Ser Ala Gly Phe Leu Pro Thr		
290	295	300
His Arg Leu Glu Pro Arg Pro Glu Pro Arg Pro Asp Pro Arg Pro Gly		
305	310	315
320		
Pro Glu Leu Pro Leu Pro Ser Ile Pro Ala Trp Thr Gly Pro Glu Ile		

325	330	335
Pro Glu Ser Gly Pro Ser Ser Gly Met Cys Gln Arg Asn Pro Gln Val		
340	345	350
Cys Gly Pro Gly Arg Cys Ile Ser Arg Pro Ser Gly Tyr Thr Cys Ala		
355	360	365
Cys Asp Ser Gly Phe Arg Leu Ser Pro Gln Gly Thr Arg Cys Ile Asp		
370	375	380
Val Asp Glu Cys Arg Arg Val Pro Pro Pro Cys Ala Pro Gly Arg Cys		
385	390	395
Glu Asn Ser Pro Gly Ser Phe Arg Cys Val Cys Gly Pro Gly Phe Arg		
405	410	415
Ala Gly Pro Arg Ala Ala Glu Cys Leu Asp Val Asp Glu Cys His Arg		
420	425	430
Val Pro Pro Pro Cys Asp Leu Gly Arg Cys Glu Asn Thr Pro Gly Ser		
435	440	445
Phe Leu Cys Val Cys Pro Ala Gly Tyr Gln Ala Ala Pro His Gly Ala		
450	455	460
Ser Cys Gln Asp Val Asp Glu Cys Thr Gln Ser Pro Gly Leu Cys Gly		
465	470	475
Arg Gly Ala Cys Lys Asn Leu Pro Gly Ser Phe Arg Cys Val Cys Pro		
485	490	495
Ala Gly Phe Arg Gly Ser Ala Cys Glu Glu Asp Val Asp Glu Cys Ala		
500	505	510
Gln Glu Pro Pro Pro Cys Gly Pro Gly Arg Cys Asp Asn Thr Ala Gly		
515	520	525
Ser Phe His Cys Ala Cys Pro Ala Gly Phe Arg Ser Arg Gly Pro Gly		
530	535	540
Ala Pro Cys Gln Asp Val Asp Glu Cys Ala Arg Ser Pro Pro Pro Cys		
545	550	555
Thr Tyr Gly Arg Cys Glu Asn Thr Glu Gly Ser Phe Gln Cys Val Cys		
565	570	575
Pro Met Gly Phe Gln Pro Asn Thr Ala Gly Ser Glu Cys Glu Asp Val		
580	585	590
Asp Glu Cys Glu Asn His Leu Ala Cys Pro Gly Gln Glu Cys Val Asn		
595	600	605
Ser Pro Gly Ser Phe Gln Cys Arg Thr Cys Pro Ser Gly His His Leu		
610	615	620
His Arg Gly Arg Cys Thr Asp Val Asp Glu Cys Ser Ser Gly Ala Pro		

625	630	635	640
Pro Cys Gly Pro His Gly His Cys Thr Asn Thr Glu Gly Ser Phe Arg			
645		650	655
Cys Ser Cys Ala Pro Gly Tyr Arg Ala Pro Ser Gly Arg Pro Gly Pro			
660	665		670
Cys Ala Asp Val Asn Glu Cys Leu Glu Gly Asp Phe Cys Phe Pro His			
675	680		685
Gly Glu Cys Leu Asn Thr Asp Gly Ser Phe Ala Cys Thr Cys Ala Pro			
690	695		700
Gly Tyr Arg Pro Gly Pro Arg Gly Ala Ser Cys Leu Asp Val Asp Glu			
705	710	715	720
Cys Ser Glu Glu Asp Leu Cys Gln Ser Gly Ile Cys Thr Asn Thr Asp			
725	730		735
Gly Ser Phe Glu Cys Ile Cys Pro Pro Gly His Arg Ala Gly Pro Asp			
740	745		750
Leu Ala Ser Cys Leu Gly Arg Gly Arg Met Ser Arg Ala Arg Pro Ser			
755	760		765
Pro Val Arg Val Ala Ala Leu			
770	775		

<210> 79
 <211> 669
 <212> PRT
 <213> Homo sapiens

<400> 79			
Arg Gly Pro Met Gly Phe Gln Pro Asn Ala Ala Gly Ser Glu Cys Glu			
1	5	10	15
Asp Val Asp Glu Cys Glu Asn His Leu Ala Cys Pro Gly Gln Glu Cys			
20	25		30
Val Asn Ser Pro Gly Ser Phe Gln Cys Arg Ala Cys Pro Ser Gly His			
35	40		45
His Leu His Arg Gly Arg Cys Thr Asp Val Asp Glu Cys Ser Ser Gly			
50	55		60
Ala Pro Pro Cys Gly Pro His Gly His Cys Thr Asn Thr Glu Gly Ser			
65	70	75	80
Phe Arg Cys Ser Cys Ala Pro Gly Tyr Arg Ala Pro Ser Gly Arg Pro			
85	90		95
Gly Pro Cys Ala Asp Val Asn Glu Cys Leu Glu Gly Asp Phe Cys Phe			
100	105		110

Pro His Gly Glu Cys Leu Asn Thr Asp Gly Ser Phe Ala Cys Thr Cys
 115 120 125
 Ala Pro Gly Tyr Arg Pro Gly Pro Arg Gly Ala Ser Cys Leu Asp Val
 130 135 140
 Asp Glu Cys Ser Glu Glu Asp Leu Cys Gln Ser Gly Ile Cys Thr Asn
 145 150 155 160
 Thr Asp Gly Ser Phe Glu Cys Ile Cys Pro Pro Gly His Arg Ala Gly
 165 170 175
 Pro Asp Leu Ala Ser Cys Leu Asp Val Asp Glu Cys Arg Glu Arg Gly
 180 185 190
 Pro Ala Leu Cys Gly Ser Gln Arg Cys Glu Asn Ser Pro Gly Ser Tyr
 195 200 205
 Arg Cys Val Arg Asp Cys Asp Pro Gly Tyr His Ala Gly Pro Glu Gly
 210 215 220
 Thr Cys Asp Asp Val Asn Glu Cys Glu Thr Leu Gln Gly Val Cys Gly
 225 230 235 240
 Ala Ala Leu Cys Glu Asn Val Glu Gly Ser Phe Leu Cys Val Cys Pro
 245 250 255
 Asn Ser Pro Glu Glu Phe Asp Pro Met Thr Gly Arg Cys Val Pro Pro
 260 265 270
 Arg Thr Ser Ala Gly Met Phe Pro Gly Ser Gln Pro Gln Ala Pro Ala
 275 280 285
 Ser Pro Val Leu Pro Ala Arg Pro Pro Pro Pro Leu Pro Arg Arg
 290 295 300
 Pro Ser Thr Pro Arg Gln Gly Pro Val Gly Ser Gly Arg Arg Glu Cys
 305 310 315 320
 Tyr Phe Asp Thr Ala Ala Pro Asp Ala Cys Asp Asn Ile Leu Ala Arg
 325 330 335
 Asn Val Thr Trp Gln Glu Cys Cys Cys Thr Val Gly Glu Gly Trp Gly
 340 345 350
 Ser Gly Cys Arg Ile Gln Gln Cys Pro Gly Thr Glu Thr Ala Glu Tyr
 355 360 365
 Gln Ser Leu Cys Pro His Gly Arg Gly Tyr Leu Ala Pro Ser Gly Asp
 370 375 380
 Leu Ser Leu Arg Arg Asp Val Asp Glu Cys Gln Leu Phe Arg Asp Gln
 385 390 395 400
 Val Cys Lys Ser Gly Val Cys Val Asn Thr Ala Pro Gly Tyr Ser Cys
 405 410 415

Tyr Cys Ser Asn Gly Tyr Tyr Tyr His Thr Gln Arg Leu Glu Cys Ile
 420 425 430
 Asp Asn Asp Glu Cys Ala Asp Glu Glu Pro Ala Cys Glu Gly Gly Arg
 435 440 445
 Cys Val Asn Thr Val Gly Ser Tyr His Cys Thr Cys Glu Pro Pro Leu
 450 455 460
 Val Leu Asp Gly Ser Gln Arg Arg Cys Val Ser Asn Glu Ser Gln Ser
 465 470 475 480
 Leu Asp Asp Asn Leu Gly Val Cys Trp Gln Glu Val Gly Ala Asp Leu
 485 490 495
 Val Cys Ser His Pro Arg Leu Asp Arg Gln Ala Thr Tyr Thr Glu Cys
 500 505 510
 Cys Cys Leu Tyr Gly Glu Ala Trp Gly Met Asp Cys Ala Leu Cys Pro
 515 520 525
 Ala Gln Asp Ser Asp Asp Phe Glu Ala Leu Cys Asn Val Leu Arg Pro
 530 535 540
 Pro Ala Tyr Ser Pro Pro Arg Pro Gly Gly Phe Gly Leu Pro Tyr Glu
 545 550 555 560
 Tyr Gly Pro Asp Leu Gly Pro Pro Tyr Gln Gly Leu Pro Tyr Gly Pro
 565 570 575
 Glu Leu Tyr Pro Pro Pro Ala Leu Pro Tyr Asp Pro Tyr Pro Pro Pro
 580 585 590
 Pro Gly Pro Phe Ala Arg Arg Glu Ala Pro Tyr Gly Ala Pro Arg Phe
 595 600 605
 Asp Met Pro Asp Phe Glu Asp Asp Gly Gly Pro Tyr Gly Glu Ser Glu
 610 615 620
 Ala Pro Ala Pro Pro Gly Pro Gly Thr Arg Trp Pro Tyr Arg Ser Arg
 625 630 635 640
 Asp Thr Arg Arg Ser Phe Pro Glu Pro Glu Glu Pro Pro Glu Gly Gly
 645 650 655
 Ser Tyr Ala Gly Ser Leu Ala Glu Pro Arg Ala Glu Phe
 660 665

<210> 80
 <211> 321
 <212> PRT
 <213> Mus musculus

<400> 80
 Met Asn Ser Thr Leu Asp Ser Ser Pro Ala Pro Gly Leu Thr Ile Ser
 1 5 10 15

Pro Thr Met Asp Leu Val Thr Trp Ile Tyr Phe Ser Val Thr Phe Leu
 20 25 30

Ala Met Ala Thr Cys Val Gly Gly Met Ala Gly Asn Ser Leu Val Ile
 35 40 45

Trp Leu Leu Ser Cys Asn Gly Met Gln Arg Ser Pro Phe Cys Val Tyr
 50 55 60

Val Leu Asn Leu Ala Val Ala Asp Phe Leu Phe Leu Phe Cys Met Ala
 65 70 75 80

Ser Met Leu Ser Leu Glu Thr Gly Pro Leu Leu Ile Val Asn Ile Ser
 85 90 95

Ala Lys Ile Tyr Glu Gly Met Arg Arg Ile Lys Tyr Phe Ala Tyr Thr
 100 105 110

Ala Gly Leu Ser Leu Leu Thr Ala Ile Ser Thr Gln Arg Cys Leu Ser
 115 120 125

Val Leu Phe Pro Ile Trp Tyr Lys Cys His Arg Pro Arg His Leu Ser
 130 135 140

Ser Val Val Ser Gly Ala Leu Trp Ala Leu Ala Phe Leu Met Asn Phe
 145 150 155 160

Leu Ala Ser Phe Phe Cys Val Gln Phe Trp His Pro Asn Lys His Gln
 165 170 175

Cys Phe Lys Val Asp Ile Val Phe Asn Ser Leu Ile Leu Gly Ile Phe
 180 185 190

Met Pro Val Met Ile Leu Thr Ser Thr Ile Leu Phe Ile Arg Val Arg
 195 200 205

Lys Asn Ser Leu Met Gln Arg Arg Arg Pro Arg Arg Leu Tyr Val Val
 210 215 220

Ile Leu Thr Ser Ile Leu Val Phe Leu Thr Cys Ser Leu Pro Leu Gly
 225 230 235 240

Ile Asn Trp Phe Leu Leu Tyr Trp Val Asp Val Lys Arg Asp Val Arg
 245 250 255

Leu Leu Tyr Ser Cys Val Ser Arg Phe Ser Ser Ser Leu Ser Ser Ser
 260 265 270

Ala Asn Pro Val Ile Tyr Phe Leu Val Gly Ser Gln Lys Ser His Arg
 275 280 285

Leu Gln Glu Ser Leu Gly Ala Val Leu Gly Arg Ala Leu Arg Asp Glu
 290 295 300

Pro Glu Pro Glu Gly Arg Glu Thr Pro Ser Thr Cys Thr Asn Asp Gly
 305 310 315 320

Val

<210> 81
<211> 322
<212> PRT
<213> Homo sapiens

<400> 81
Met Asp Pro Thr Ile Ser Thr Leu Asp Thr Glu Leu Thr Pro Ile Asn
1 5 10 15
Gly Thr Glu Glu Thr Leu Cys Tyr Lys Gln Thr Leu Ser Leu Thr Val
20 25 30
Leu Thr Cys Ile Val Ser Leu Val Gly Leu Thr Gly Asn Ala Val Val
35 40 45
Leu Trp Leu Leu Gly Cys Arg Met Arg Arg Asn Ala Phe Ser Ile Tyr
50 55 60
Ile Leu Asn Leu Ala Ala Ala Asp Phe Leu Phe Leu Ser Gly Arg Leu
65 70 75 80
Ile Tyr Ser Leu Leu Ser Phe Ile Ser Ile Pro His Thr Ile Ser Lys
85 90 95
Ile Leu Tyr Pro Val Met Met Phe Ser Tyr Phe Ala Gly Leu Ser Phe
100 105 110
Leu Ser Ala Val Ser Thr Glu Arg Cys Leu Ser Val Leu Trp Pro Ile
115 120 125
Trp Tyr Arg Cys His Arg Pro Thr His Leu Ser Ala Val Val Cys Val
130 135 140
Leu Leu Trp Ala Leu Ser Leu Leu Arg Ser Ile Leu Glu Trp Met Leu
145 150 155 160
Cys Gly Phe Leu Phe Ser Gly Ala Asp Ser Ala Trp Cys Gln Thr Ser
165 170 175
Asp Phe Ile Thr Val Ala Trp Leu Ile Phe Leu Cys Val Val Leu Cys
180 185 190
Gly Ser Ser Leu Val Leu Leu Ile Arg Ile Leu Cys Gly Ser Arg Lys
195 200 205
Ile Pro Leu Thr Arg Leu Tyr Val Thr Ile Leu Leu Thr Val Leu Val
210 215 220
Phe Leu Leu Cys Gly Leu Pro Phe Gly Ile Gln Phe Phe Leu Phe Leu
225 230 235 240
Trp Ile His Val Asp Arg Glu Val Leu Phe Cys His Val His Leu Val

245	250	255
Ser Ile Phe Leu Ser Ala Leu Asn Ser Ser Ala Asn Pro Ile Ile Tyr		
260	265	270
Phe Phe Val Gly Ser Phe Arg Gln Arg Gln Asn Arg Gln Asn Leu Lys		
275	280	285
Leu Val Leu Gln Arg Ala Leu Gln Asp Ala Ser Glu Val Asp Glu Gly		
290	295	300
Gly Gly Gln Leu Pro Glu Glu Ile Leu Glu Leu Ser Gly Ser Arg Leu		
305	310	315
Glu Gln		

<210> 82
<211> 304
<212> PRT
<213> Homo sapiens

<400> 82		
Met Asp Asn Thr Ile Pro Gly Gly Ile Asn Ile Thr Ile Leu Ile Pro		
1	5	10
Asn Leu Met Ile Ile Ile Phe Gly Leu Val Gly Leu Thr Gly Asn Gly		
20	25	30
Ile Val Phe Trp Leu Leu Gly Phe Cys Leu His Arg Asn Ala Phe Ser		
35	40	45
Val Tyr Ile Leu Asn Leu Ala Leu Asp Phe Phe Leu Leu Gly		
50	55	60
His Ile Ile Asp Ser Ile Leu Leu Leu Asn Val Phe Tyr Pro Ile		
65	70	75
Thr Phe Leu Leu Cys Phe Tyr Thr Ile Met Met Val Leu Tyr Ile Ala		
85	90	95
Gly Leu Ser Met Leu Ser Ala Ile Ser Thr Glu Arg Cys Leu Ser Val		
100	105	110
Leu Cys Pro Ile Trp Tyr His Cys His Arg Pro Glu His Thr Ser Thr		
115	120	125
Val Met Cys Ala Val Ile Trp Val Leu Ser Leu Leu Ile Cys Ile Leu		
130	135	140
Asn Ser Tyr Phe Cys Gly Phe Leu Asn Thr Gln Tyr Lys Asn Glu Asn		
145	150	155
Gly Cys Leu Ala Leu Asn Phe Phe Thr Ala Ala Tyr Leu Met Phe Leu		
165	170	175

Phe	Val	Val	Leu	Cys	Leu	Ser	Ser	Leu	Ala	Leu	Val	Ala	Arg	Leu	Phe
180								185						190	
Cys	Gly	Thr	Gly	Gln	Ile	Lys	Leu	Thr	Arg	Leu	Tyr	Val	Thr	Ile	Ile
195							200						205		
Leu	Ser	Ile	Leu	Val	Phe	Leu	Leu	Cys	Gly	Leu	Pro	Phe	Gly	Ile	His
210					215							220			
Trp	Phe	Leu	Leu	Phe	Lys	Ile	Lys	Asp	Asp	Phe	His	Val	Phe	Asp	Leu
225					230				235					240	
Gly	Phe	Tyr	Leu	Ala	Ser	Val	Val	Leu	Thr	Ala	Ile	Asn	Ser	Cys	Ala
245							250						255		
Asn	Pro	Ile	Ile	Tyr	Phe	Phe	Val	Gly	Ser	Phe	Arg	His	Arg	Leu	Lys
260							265						270		
His	Gln	Thr	Leu	Lys	Met	Val	Leu	Gln	Asn	Ala	Leu	Gln	Asp	Thr	Pro
275						280							285		
Glu	Thr	Ala	Lys	Ile	Met	Val	Glu	Met	Ser	Arg	Ser	Lys	Ser	Glu	Pro
290					295							300			

<210> 83
<211> 321
<212> PRT
<213> Homo sapiens

<400> 83															
Met	Glu	Pro	Leu	Ala	Met	Thr	Leu	Tyr	Pro	Leu	Glu	Ser	Thr	Gln	Pro
1					5					10					15
Thr	Arg	Asn	Lys	Thr	Pro	Asn	Glu	Thr	Thr	Trp	Ser	Ser	Glu	His	Thr
20							25						30		
Asp	Asp	His	Thr	Tyr	Phe	Leu	Val	Ser	Leu	Val	Ile	Cys	Ser	Leu	Gly
35						40						45			
Leu	Ala	Gly	Asn	Gly	Leu	Leu	Ile	Trp	Phe	Leu	Ile	Phe	Cys	Ile	Lys
50					55						60				
Arg	Lys	Pro	Phe	Thr	Ile	Tyr	Ile	Leu	His	Leu	Ala	Ile	Ala	Asp	Phe
65					70				75					80	
Met	Val	Leu	Leu	Cys	Ser	Ser	Ile	Met	Lys	Leu	Val	Asn	Thr	Phe	His
85							90						95		
Ile	Tyr	Asn	Met	Thr	Leu	Glu	Ser	Tyr	Ala	Ile	Leu	Phe	Met	Ile	Phe
100						105						110			
Gly	Tyr	Asn	Thr	Gly	Leu	His	Leu	Leu	Thr	Ala	Ile	Ser	Val	Glu	Arg
115						120						125			

Cys Leu Ser Val Leu Tyr Pro Ile Trp Tyr Gln Cys Gln Arg Pro Lys
 130 135 140
 His Gln Ser Ala Val Ala Cys Met Leu Leu Trp Ala Leu Ser Val Leu
 145 150 155 160
 Val Ser Gly Leu Glu Asn Phe Phe Cys Ile Leu Glu Val Lys Pro Gln
 165 170 175
 Phe Pro Glu Cys Arg Tyr Val Tyr Ile Phe Ser Cys Ile Leu Thr Phe
 180 185 190
 Leu Val Phe Val Pro Leu Met Ile Phe Ser Asn Leu Ile Leu Phe Ile
 195 200 205
 Gln Val Cys Cys Asn Leu Lys Pro Arg Gln Pro Thr Lys Leu Tyr Val
 210 215 220
 Ile Ile Met Thr Thr Val Ile Leu Phe Leu Val Phe Ala Met Pro Met
 225 230 235 240
 Lys Val Leu Leu Ile Ile Gly Tyr Tyr Ser Ser Ser Leu Asp Asp Ser
 245 250 255
 Val Trp Asp Ser Leu Pro Tyr Leu Asn Met Leu Ser Thr Ile Asn Cys
 260 265 270
 Ser Ile Asn Pro Ile Val Tyr Phe Val Val Gly Ser Leu Arg Arg Lys
 275 280 285
 Arg Ser Arg Lys Ser Leu Lys Glu Ala Leu Gln Lys Val Phe Glu Glu
 290 295 300
 Lys Pro Val Val Ala Ser Arg Glu Asn Val Thr Gln Phe Ser Leu Pro
 305 310 315 320
 Ser

<210> 84
 <211> 322
 <212> PRT
 <213> Homo sapiens

<400> 84
 Met Asp Pro Thr Val Pro Val Phe Gly Thr Lys Leu Thr Pro Ile Asn
 1 5 10 15
 Gly Arg Glu Glu Thr Pro Cys Tyr Asn Gln Thr Leu Ser Phe Thr Val
 20 25 30
 Leu Thr Cys Ile Ile Ser Leu Val Gly Leu Thr Gly Asn Ala Val Val
 35 40 45
 Leu Trp Leu Leu Gly Tyr Arg Met Arg Arg Asn Ala Val Ser Ile Tyr

50	55	60
Ile Leu Asn Leu Ala Ala Asp Phe Leu Phe Leu Ser Phe Gln Ile		
65	70	75
Ile Arg Ser Pro Leu Arg Leu Ile Asn Ile Ser His Leu Ile Arg Lys		
85	90	95
Ile Leu Val Ser Val Met Thr Phe Pro Tyr Phe Thr Gly Leu Ser Met		
100	105	110
Leu Ser Ala Ile Ser Thr Glu Arg Cys Leu Ser Val Leu Trp Pro Ile		
115	120	125
Trp Tyr Arg Cys Arg Arg Pro Thr His Leu Ser Ala Val Val Cys Val		
130	135	140
Leu Leu Trp Gly Leu Ser Leu Leu Phe Ser Met Leu Glu Trp Arg Phe		
145	150	155
Cys Asp Phe Leu Phe Ser Gly Ala Asp Ser Ser Trp Cys Glu Thr Ser		
165	170	175
Asp Phe Ile Pro Val Ala Trp Leu Ile Phe Leu Cys Val Val Leu Cys		
180	185	190
Val Ser Ser Leu Val Leu Leu Val Arg Ile Leu Cys Gly Ser Arg Lys		
195	200	205
Met Pro Leu Thr Arg Leu Tyr Val Thr Ile Leu Leu Thr Val Leu Val		
210	215	220
Phe Leu Leu Cys Gly Leu Pro Phe Gly Ile Leu Gly Ala Leu Ile Tyr		
225	230	235
Arg Met His Leu Asn Leu Glu Val Leu Tyr Cys His Val Tyr Leu Val		
245	250	255
Cys Met Ser Leu Ser Ser Leu Asn Ser Ser Ala Asn Pro Ile Ile Tyr		
260	265	270
Phe Phe Val Gly Ser Phe Arg Gln Arg Gln Asn Arg Gln Asn Leu Lys		
275	280	285
Leu Val Leu Gln Arg Ala Leu Gln Asp Lys Pro Glu Val Asp Lys Gly		
290	295	300
Glu Gly Gln Leu Pro Glu Glu Ser Leu Glu Leu Ser Gly Ser Arg Leu		
305	310	315
Gly Pro		

<210> 85
<211> 149
<212> PRT

<213> Mus pahari

<400> 85
Met Gly Leu Glu Lys Ser Leu Ile Leu Phe Pro Leu Phe Val Leu Leu
1 5 10 15

Leu Gly Trp Val Gln Pro Ser Leu Gly Lys Glu Ser Ser Ala Gln Lys
20 25 30

Phe Glu Arg Gln His Met Asp Ser Ser Gly Ser Ser Asn Asn Ser Pro
35 40 45

Thr Tyr Cys Asn Gln Met Met Lys Ser Arg Ser Met Thr Lys Glu Ser
50 55 60

Cys Lys Pro Val Asn Thr Phe Val His Glu Pro Leu Glu Asp Val Gln
65 70 75 80

Ala Ile Cys Ser Gln Glu Asn Val Thr Cys Lys Asn Gly Asn Arg Asn
85 90 95

Cys Tyr Lys Ser Ser Ser Ala Leu His Ile Thr Asp Cys His Leu Lys
100 105 110

Gly Asn Ser Lys Tyr Pro Asn Cys Asn Tyr Asn Thr Asn Gln Tyr Gln
115 120 125

Lys His Ile Ile Val Ala Cys Asp Gly Asn Pro Tyr Val Pro Val His
130 135 140

Leu Asp Ala Thr Val
145

<210> 86
<211> 129
<212> PRT
<213> Homo sapiens

<220>
<221> VARIANT
<222> (1)
<223> Wherein Xaa is any amino acid as defined in the
specification

<400> 86
Xaa Lys Glu Ser Ala Ala Ala Lys Phe Glu Arg Gln His Met Asp Ser
1 5 10 15

Gly Asn Ser Pro Ser Ser Ser Thr Tyr Cys Asn Gln Met Met Arg
20 25 30

Arg Arg Asn Met Thr Gln Gly Arg Cys Lys Pro Val Asn Thr Phe Val
35 40 45

His Glu Ser Leu Val Asp Val Gln Asn Val Cys Phe Gln Glu Lys Val
50 55 60

Thr Cys Lys Asn Gly Gln Gly Asn Cys Tyr Lys Ser Asn Ser Ser Met
65 70 75 80

His Ile Thr Asp Cys Arg Leu Thr Asn Gly Ser Arg Tyr Pro Asn Cys
85 90 95

Ala Tyr Arg Thr Ser Pro Lys Glu Arg His Ile Ile Val Ala Cys Glu
100 105 110

Gly Ser Pro Tyr Val Pro Val His Phe Asp Ala Ser Val Glu Asp Ser
115 120 125

Thr

<210> 87
<211> 208
<212> PRT
<213> Mus musculus

<400> 87
Met Lys Val Thr Leu Val His Leu Leu Phe Met Met Leu Leu Leu Leu
1 5 10 15

Leu Gly Leu Gly Leu Gly Leu Gly Leu His Met Ala Ala Ala
20 25 30

Val Leu Glu Asp Gln Pro Leu Asn Glu Phe Trp Pro Ser Asp Ser Gln
35 40 45

Asn Thr Glu Glu Gly Glu Ile Trp Thr Thr Glu Gly Leu Ala Leu
50 55 60

Gly Tyr Lys Glu Met Ala Gln Pro Val Trp Pro Glu Glu Ala Val Leu
65 70 75 80

Ser Glu Asp Glu Val Gly Ser Arg Met Leu Arg Ala Glu Pro Arg
85 90 95

Phe Gln Ser Lys Gln Asp Tyr Leu Lys Phe Asp Leu Ser Val Arg Asp
100 105 110

Cys Asn Thr Met Met Ala His Lys Ile Lys Glu Pro Asn Gln Ser Cys
115 120 125

Ile Asn Gln Tyr Thr Phe Ile His Glu Asp Pro Asn Thr Val Lys Ala
130 135 140

Val Cys Asn Gly Ser Leu Val Asp Cys Asp Leu Gln Gly Lys Cys
145 150 155 160

Tyr Lys Ser Pro Arg Pro Phe Asp Leu Thr Leu Cys Lys Leu Ala Lys
165 170 175

Pro Gly Gln Val Thr Pro Asn Cys His Tyr Leu Thr Tyr Ile Thr Glu

180

185

190

Lys Ser Ile Phe Met Thr Cys Asn Asp Lys Arg Gln Leu Glu Thr Lys
195 200 205

<210> 88

<211> 128

<212> PRT

<213> Presbytis entellus

<400> 88

Gly Glu Ser Arg Ala Glu Lys Phe Gln Arg Gln His Met Asp Ser Gly
1 5 10 15

Ser Ser Pro Ser Ser Ser Thr Tyr Cys Asn Gln Met Met Lys Leu
20 25 30

Arg Asn Met Thr Gln Gly Ser Cys Lys Ser Val Asn Thr Phe Val His
35 40 45

Glu Pro Leu Val Asp Val Gln Asn Val Cys Phe Gln Glu Lys Val Thr
50 55 60

Cys Lys Asn Gly Gln Thr Asn Cys Phe Lys Ser Asn Ser Arg Met His
65 70 75 80

Ile Thr Glu Cys Arg Leu Thr Asn Gly Ser Lys Tyr Pro Asn Cys Ala
85 90 95

Tyr Gly Thr Ser Pro Lys Glu Arg His Ile Ile Val Ala Cys Glu Gly
100 105 110

Ser Pro Tyr Val Pro Val His Phe Asp Asp Ser Val Glu Asp Ser Thr
115 120 125

<210> 89

<211> 119

<212> PRT

<213> Iguana iguana

<400> 89

Gln Asp Trp Ser Ser Phe Gln Asn Lys His Ile Asp Tyr Pro Glu Thr
1 5 10 15

Ser Ala Ser Asn Pro Asn Ala Tyr Cys Asp Leu Met Met Gln Arg Arg
20 25 30

Asn Leu Asn Pro Thr Lys Cys Lys Thr Arg Asn Thr Phe Val His Ala
35 40 45

Ser Pro Ser Glu Ile Gln Gln Val Cys Gly Ser Gly Gly Thr His Tyr
 50 55 60

Glu Asp Asn Leu Tyr Asp Ser Asn Glu Ser Phe Asp Leu Thr Asp Cys
 65 70 75 80

Lys Asn Val Gly Gly Thr Ala Pro Ser Ser Cys Lys Tyr Asn Gly Thr
 85 90 95

Pro Gly Thr Lys Arg Ile Arg Ile Ala Cys Glu Asn Asn Gln Pro Val
 100 105 110

His Phe Glu Leu Val Leu Ser
 115

`
 <210> 90
 <211> 493
 <212> PRT
 <213> Homo sapiens

<400> 90
 Met Cys Glu Leu Tyr Ser Lys Arg Asp Thr Leu Gly Leu Arg Lys Lys
 1 5 10 15

His Ile Gly Pro Ser Cys Lys Val Phe Phe Ala Ser Asp Pro Ile Lys
 20 25 30

Ile Val Arg Ala Gln Arg Gln Tyr Met Phe Asp Glu Asn Gly Glu Gln
 35 40 45

Tyr Leu Asp Cys Ile Asn Asn Val Ala His Gly Val Val Lys Ala Ala
 50 55 60

Leu Lys Gln Met Glu Leu Leu Asn Thr Asn Ser Arg Phe Leu His Asp
 65 70 75 80

Asn Ile Val Glu Tyr Ala Lys Arg Leu Ser Ala Thr Leu Pro Glu Lys
 85 90 95

Leu Ser Val Cys Tyr Phe Thr Asn Ser Gly Ser Glu Ala Asn Asp Leu
 100 105 110

Ala Leu Arg Leu Ala Arg Gln Phe Arg Gly His Gln Asp Val Ile Thr
 115 120 125

Leu Asp His Ala Tyr His Gly His Leu Ser Ser Leu Ile Glu Ile Ser
 130 135 140

Pro Tyr Lys Phe Gln Lys Gly Lys Asp Val Lys Lys Glu Phe Val His
 145 150 155 160

Val Ala Pro Thr Pro Asp Thr Tyr Arg Gly Lys Tyr Arg Glu Asp His
 165 170 175

Ala Asp Ser Ala Ser Ala Tyr Ala Asp Glu Val Lys Lys Ile Ile Glu

180	185	190
Asp Ala His Asn Ser Gly Arg Lys Ile Ala Ala Phe Ile Ala Glu Ser		
195	200	205
Met Gln Ser Cys Gly Gly Gln Ile Ile Pro Pro Ala Gly Tyr Phe Gln		
210	215	220
Lys Val Ala Glu Tyr Val His Gly Ala Gly Gly Val Phe Ile Ala Asp		
225	230	235
Glu Val Gln Val Gly Phe Gly Arg Val Gly Lys His Phe Trp Ser Phe		
245	250	255
Gln Met Tyr Gly Glu Asp Phe Val Pro Asp Ile Val Thr Met Gly Lys		
260	265	270
Pro Met Gly Asn Gly His Pro Val Ala Cys Val Val Thr Thr Lys Glu		
275	280	285
Ile Ala Glu Ala Phe Ser Ser Ser Gly Met Glu Tyr Phe Asn Thr Tyr		
290	295	300
Gly Gly Asn Pro Val Ser Cys Ala Val Gly Leu Ala Val Leu Asp Ile		
305	310	315
Ile Glu Asn Glu Asp Leu Gln Gly Asn Ala Lys Arg Val Gly Asn Tyr		
325	330	335
Leu Thr Glu Leu Leu Lys Lys Gln Lys Ala Lys His Thr Leu Ile Gly		
340	345	350
Asp Ile Arg Gly Ile Gly Leu Phe Ile Gly Ile Asp Leu Val Lys Asp		
355	360	365
His Leu Lys Arg Thr Pro Ala Thr Ala Glu Ala Gln His Ile Ile Tyr		
370	375	380
Lys Met Lys Glu Lys Arg Val Leu Leu Ser Ala Asp Gly Pro His Arg		
385	390	395
Asn Val Leu Lys Ile Lys Pro Pro Met Cys Phe Thr Glu Glu Asp Ala		
405	410	415
Lys Phe Met Val Asp Gln Leu Asp Arg Ile Leu Thr Val Leu Glu Glu		
420	425	430
Ala Met Gly Thr Lys Thr Glu Ser Val Thr Ser Glu Asn Thr Pro Cys		
435	440	445
Lys Thr Lys Met Leu Lys Glu Ala His Ile Glu Leu Leu Arg Asp Ser		
450	455	460
Thr Thr Asp Ser Lys Glu Asn Pro Ser Arg Lys Arg Asn Gly Met Cys		
465	470	475
Thr Asp Thr His Ser Leu Leu Ser Lys Arg Leu Lys Thr		

485

490

<210> 91
<211> 499
<212> PRT
<213> Mus musculus

<400> 91
Met Cys Glu Leu Tyr Ser Lys Gln Asp Thr Leu Ala Leu Arg Glu Arg
1 5 10 15
His Ile Gly Pro Ser Cys Lys Ile Phe Phe Ala Ala Asp Pro Ile Lys
20 25 30
Ile Met Arg Ala Gln Gly Gln Tyr Met Phe Asp Glu Lys Gly Glu Arg
35 40 45
Tyr Leu Asp Cys Ile Asn Asn Val Ala His Val Gly His Cys His Pro
50 55 60
Glu Val Val Lys Ala Ala Ala Lys Gln Met Glu Leu Leu Asn Thr Asn
65 70 75 80
Ser Arg Phe Leu His Asp Asn Ile Ile Glu Phe Ala Lys Arg Leu Thr
85 90 95
Ala Thr Leu Pro Gln Glu Leu Ser Val Cys Tyr Phe Thr Asn Ser Gly
100 105 110
Ser Glu Ala Asn Asp Leu Ala Leu Arg Leu Ala Arg Gln Phe Arg Gly
115 120 125
His Gln Asp Val Ile Thr Leu Asp His Ala Tyr His Gly His Leu Ser
130 135 140
Ser Leu Ile Glu Ile Ser Pro Tyr Lys Phe Gln Lys Gly Lys Asp Val
145 150 155 160
Lys Arg Glu Thr Val His Val Ala Pro Ala Pro Asp Thr Tyr Arg Gly
165 170 175
Lys Tyr Arg Glu Asp His Glu Asp Pro Ser Thr Ala Tyr Ala Asp Glu
180 185 190
Val Lys Lys Ile Ile Glu Glu Ala His Ser Ser Gly Arg Lys Ile Ala
195 200 205
Ala Phe Ile Ala Glu Ser Met Gln Ser Cys Gly Gly Gln Ile Ile Pro
210 215 220
Pro Ala Gly Tyr Phe Gln Lys Val Ala Glu His Ile His Lys Ala Gly
225 230 235 240
Gly Val Phe Ile Ala Asp Glu Val Gln Val Gly Phe Gly Arg Val Gly
245 250 255

Arg Tyr Phe Trp Ser Phe Gln Met Tyr Gly Glu Asp Phe Val Pro Asp
 260 265 270
 Ile Val Thr Met Gly Lys Pro Met Gly Asp Gly His Pro Ile Ser Cys
 275 280 285
 Val Val Thr Thr Lys Glu Ile Ala Glu Ala Phe Ser Ser Ser Gly Met
 290 295 300
 Glu Tyr Phe Asn Thr Tyr Gly Gly Asn Pro Val Ser Cys Ala Val Gly
 305 310 315 320
 Leu Ala Val Leu Asp Val Ile Glu Lys Glu Asn Leu Gln Gly Asn Ala
 325 330 335
 Val Arg Val Gly Thr Tyr Leu Met Glu Leu Leu Ser Glu Gln Lys Ala
 340 345 350
 Lys His Pro Leu Ile Gly Asp Ile Arg Gly Val Gly Leu Phe Ile Gly
 355 360 365
 Ile Asp Leu Val Lys Asp Arg Glu Lys Arg Thr Pro Ala Thr Ala Glu
 370 375 380
 Ala Gln His Ile Ile Tyr Glu Met Lys Gly Lys Gly Val Leu Leu Ser
 385 390 395 400
 Ala Asp Gly Pro His Arg Asn Val Leu Lys Ile Lys Pro Pro Met Cys
 405 410 415
 Phe Thr Glu Asp Asp Ala Lys Phe Leu Val Asp His Leu Asp Gly Ile
 420 425 430
 Leu Thr Val Leu Glu Glu Ala Met Asp Ser Lys Ser Gly Thr Val Phe
 435 440 445
 Ser Glu Asn Thr Ala Tyr Arg Thr Lys Met Pro Lys Glu Ile Gln Val
 450 455 460
 Glu Leu Pro Asn Leu Ser Ala Thr Glu Ala Arg Glu Ile Pro Arg Gly
 465 470 475 480
 Lys Arg Asn Gly Val Cys Ser Asp Gln Gln Ala Leu Leu Ser Lys Arg
 485 490 495
 Leu Lys Thr

<210> 92
 <211> 426
 <212> PRT
 <213> Homo sapiens

<400> 92
 Met Glu Leu Leu Asn Thr Asn Ser Arg Phe Leu His Asp Asn Ile Val
 1 5 10 15

Glu Tyr Ala Lys Arg Leu Ser Ala Thr Leu Pro Glu Lys Leu Ser Val
 20 25 30

 Cys Tyr Phe Thr Asn Ser Gly Ser Glu Ala Asn Asp Leu Ala Leu Arg
 35 40 45

 Leu Ala Arg Gln Phe Arg Gly His Gln Asp Val Ile Thr Leu Asp His
 50 55 60

 Ala Tyr His Gly His Leu Ser Ser Leu Ile Glu Ile Ser Pro Tyr Lys
 65 70 75 80

 Phe Gln Lys Gly Lys Asp Val Lys Lys Glu Phe Val His Val Ala Pro
 85 90 95

 Thr Pro Asp Thr Tyr Arg Gly Lys Tyr Arg Glu Asp His Ala Asp Ser
 100 105 110

 Ala Ser Ala Tyr Ala Asp Glu Val Lys Lys Ile Ile Glu Asp Ala His
 115 120 125

 Asn Ser Gly Arg Lys Ile Ala Ala Phe Ile Ala Glu Ser Met Gln Ser
 130 135 140

 Cys Gly Gly Gln Ile Ile Pro Pro Ala Gly Tyr Phe Gln Lys Val Ala
 145 150 155 160

 Glu Tyr Val His Gly Ala Gly Val Phe Ile Ala Asp Glu Val Gln
 165 170 175

 Val Gly Phe Gly Arg Val Gly Lys His Phe Trp Ser Phe Gln Met Tyr
 180 185 190

 Gly Glu Asp Phe Val Pro Asp Ile Val Thr Met Gly Lys Pro Met Gly
 195 200 205

 Asn Gly His Pro Val Ala Cys Val Val Thr Thr Lys Glu Ile Ala Glu
 210 215 220

 Ala Phe Ser Ser Ser Gly Met Glu Tyr Phe Asn Thr Tyr Gly Gly Asn
 225 230 235 240

 Pro Val Ser Cys Ala Val Gly Leu Ala Val Leu Asp Ile Ile Glu Asn
 245 250 255

 Glu Asp Leu Gln Gly Asn Ala Lys Arg Val Gly Asn Tyr Leu Thr Glu
 260 265 270

 Leu Leu Lys Lys Gln Lys Ala Lys His Thr Leu Ile Gly Asp Ile Arg
 275 280 285

 Gly Ile Gly Leu Phe Ile Gly Ile Asp Leu Val Lys Asp His Leu Lys
 290 295 300

 Arg Thr Pro Ala Thr Ala Glu Ala Gln His Ile Ile Tyr Lys Met Lys
 305 310 315 320

Glu Lys Arg Val Leu Leu Ser Ala Asp Gly Pro His Arg Asn Val Leu
325 330 335

Lys Ile Lys Pro Pro Met Cys Phe Thr Glu Glu Asp Ala Lys Phe Met
340 345 350

Val Asp Gln Leu Asp Arg Ile Leu Thr Val Leu Glu Glu Ala Met Gly
355 360 365

Thr Lys Thr Glu Ser Val Thr Ser Glu Asn Thr Pro Cys Lys Thr Lys
370 375 380

Met Leu Lys Glu Ala His Ile Glu Leu Leu Arg Asp Ser Thr Thr Asp
385 390 395 400

Ser Lys Glu Asn Pro Ser Arg Lys Arg Asn Gly Met Cys Thr Asp Thr
405 410 415

His Ser Leu Leu Ser Lys Arg Leu Lys Thr
420 425

<210> 93

<211> 473

<212> PRT

<213> Mus musculus

<400> 93

Thr Arg Thr Ala Arg Arg His Gly Arg Gly His Gly Ala Lys Ala Val
1 5 10 15

Thr Leu Asp Leu Arg Arg Leu Leu Ser Ser Ser Cys Arg Leu Phe
20 25 30

Phe Pro Glu Asp Pro Val Lys Ile Ile Arg Gly Gln Gly Gln Tyr Leu
35 40 45

Tyr Asp Glu Gln Gly Arg Glu Tyr Leu Asp Cys Ile Asn Asn Val Ala
50 55 60

His Val Gly His Cys His Pro Thr Val Val Gln Ala Ala His Glu Gln
65 70 75 80

Asn Leu Val Leu Asn Thr Asn Ser Arg Tyr Leu His Gly Asn Ile Val
85 90 95

Asp Tyr Ala Gln Arg Leu Ser Glu Thr Leu Pro Glu Gln Leu Ser Val
100 105 110

Phe Tyr Phe Leu Asn Ser Gly Ser Glu Ala Asn Asp Leu Ala Leu Arg
115 120 125

Leu Ala Arg Gln Tyr Thr Gly His Gln Asp Val Val Leu Asp His
130 135 140

Ala Tyr His Gly His Leu Ser Ser Leu Ile Asp Ile Ser Pro Tyr Lys

145	150	155	160
Phe Arg Asn Leu Gly Gly Gln Lys Glu Trp Val His Val Ala Pro Leu			
165	170	175	
Pro Asp Thr Tyr Arg Gly Pro Tyr Arg Glu Asp His Pro Asn Pro Ala			
180	185	190	
Glu Ala Tyr Ala Asn Glu Val Lys His Val Ile Ser Ser Ala Gln Gln			
195	200	205	
Lys Gly Arg Lys Ile Ala Ala Phe Phe Ala Glu Ser Leu Pro Ser Val			
210	215	220	
Ser Gly Gln Ile Ile Pro Pro Ala Gly Tyr Phe Ser Gln Val Ala Glu			
225	230	235	240
His Ile His Arg Ala Gly Gly Leu Phe Val Ala Asp Glu Ile Gln Val			
245	250	255	
Gly Phe Gly Arg Ile Gly Lys His Phe Trp Ala Phe Gln Leu Glu Gly			
260	265	270	
Glu Asp Phe Val Pro Asp Ile Val Thr Met Gly Lys Ser Ile Gly Asn			
275	280	285	
Gly His Pro Val Ala Cys Met Ala Thr Thr Gln Ala Val Ser Arg Ala			
290	295	300	
Phe Glu Ala Thr Gly Val Glu Tyr Phe Asn Thr Phe Gly Gly Asn Pro			
305	310	315	320
Val Ser Cys Ala Val Gly Leu Ala Val Leu Asp Val Leu Lys Thr Glu			
325	330	335	
Gln Leu Gln Ala His Ala Thr Asn Val Gly Ser Phe Leu Leu Glu His			
340	345	350	
Leu Thr Gln Gln Lys Ala Lys His Pro Ile Ile Gly Asp Val Arg Gly			
355	360	365	
Thr Gly Leu Phe Ile Gly Val Asp Leu Ile Lys Asp Glu Thr Leu Arg			
370	375	380	
Thr Pro Ala Thr Glu Glu Ala Glu Tyr Leu Val Ser Arg Leu Lys Glu			
385	390	395	400
Asn Tyr Ile Leu Leu Ser Ile Asp Gly Pro Gly Lys Asn Ile Leu Lys			
405	410	415	
Phe Lys Pro Pro Met Cys Phe Asn Val Asp Asn Ala Gln His Val Val			
420	425	430	
Ala Lys Leu Asp Asp Ile Leu Thr Asp Met Glu Glu Lys Val Arg Ser			
435	440	445	
Cys Glu Thr Leu Arg Ile Lys His Pro Pro Glu Asp Thr His Pro Thr			

450

455

460

Gln Ile Leu Leu Thr Arg Gln Gln Asp
 465 470

<210> 94
 <211> 494
 <212> PRT
 <213> Drosophila melanogaster

<400> 94
 Met Pro Phe Ala His Glu Gln Leu Asn Leu Val Ala Ser Glu Gln Leu
 1 5 10 15

Ser Lys Thr Glu Thr Ile Lys Leu Arg Asn Gln His Ile Gly Gln Ala
 20 25 30

Cys Gln Leu Phe Tyr Arg Ser Asp Pro Leu Lys Ile Val Arg Gly Gln
 35 40 45

Gly Gln Tyr Met Phe Asp Glu Glu Gly Thr Arg Tyr Leu Asp Cys Ile
 50 55 60

Asn Asn Val Ala His Val Gly His Cys His Pro Glu Val Val Arg Ala
 65 70 75 80

Gly Ala Leu Gln Met Ala Thr Ile Ser Thr Asn Asn Arg Phe Leu His
 85 90 95

Asp Glu Leu Val Gln Cys Ala Arg Thr Leu Thr Ser Lys Met Pro Glu
 100 105 110

Pro Leu Ser Val Cys Phe Phe Val Asn Ser Gly Ser Glu Ala Asn Asp
 115 120 125

Leu Ala Leu Arg Leu Ala Arg Asn Phe Thr Lys Arg Gln Asp Val Ile
 130 135 140

Thr Leu Asp His Ala Tyr His Gly His Leu Gln Ser Val Met Glu Val
 145 150 155 160

Ser Pro Tyr Lys Phe Asn Gln Pro Gly Gly Glu Ala Lys Pro Asp Tyr
 165 170 175

Val His Val Ala Pro Cys Pro Asp Val Tyr Gly Gly Lys Phe Thr Asp
 180 185 190

Lys Met Tyr Pro Asp Ala Asp Met Gly Ala Leu Tyr Ala Gln Pro Ile
 195 200 205

Glu Glu Ile Cys Gln Lys Gln Leu Ala Lys Gly Gln Gly Val Ala Ala
 210 215 220

Phe Ile Ala Glu Ser Leu Gln Ser Cys Gly Gly Gln Ile Leu Pro Pro
 225 230 235 240

Ala Gly Tyr Phe Gln Ala Val Tyr Asp Ala Val Arg Ser Ala Gly Gly
 245 250 255
 Val Cys Ile Ala Asp Glu Val Gln Val Gly Phe Gly Arg Val Gly Ser
 260 265 270
 His Tyr Trp Ala Phe Glu Thr Gln Asn Val Ile Pro Asp Ile Val Cys
 275 280 285
 Val Ala Lys Pro Met Gly Asn Gly His Pro Val Gly Ala Val Val Thr
 290 295 300
 Thr Pro Glu Ile Ala Gln Ala Phe His Ala Thr Gly Val Ala Tyr Phe
 305 310 315 320
 Asn Thr Tyr Gly Gly Asn Pro Val Ser Cys Ala Ile Ala Asn Ala Val
 325 330 335
 Met Arg Val Ile Glu Glu Glu Gly Leu Gln Gln Lys Ala Leu Val Leu
 340 345 350
 Gly Asp Tyr Leu Leu Glu Glu Cys Asn Arg Leu Lys Gln Glu Phe Glu
 355 360 365
 Cys Ile Gly Asp Val Arg Gly Ala Gly Leu Phe Val Gly Ile Glu Leu
 370 375 380
 Val Gln Asp Arg Lys Glu Arg Ile Pro Asp Lys Lys Ala Ala His Trp
 385 390 395 400
 Val Val Asn Arg Met Lys Gln Leu His Arg Val Leu Val Ser Ser Asp
 405 410 415
 Gly Pro Asn Asp Asn Val Ile Lys Leu Lys Pro Pro Met Cys Phe Asn
 420 425 430
 Arg Glu Asn Ala Asp Glu Phe Leu Leu Gly Phe Arg Glu Cys Leu Thr
 435 440 445
 Ala Val Met Gln Glu Arg Leu Ala Ser Ala Thr Ser Ala Ala Met Ala
 450 455 460
 Ala Thr Ser Gly Val Ile Ala Thr Ala Thr Glu Thr Leu Ala Asn Lys
 465 470 475 480
 Thr Lys Leu Phe Glu Arg Gln Asp Arg Leu Ile Lys Ser Val
 485 490

<210> 95
 <211> 1013
 <212> PRT
 <213> Mus musculus

<400> 95
 Met Gly Leu Gln Ala Leu Ser Pro Arg Met Leu Leu Trp Leu Val Val
 1 5 10 15

Ser Gly Ile Val Phe Ser Arg Val Leu Trp Val Cys Ala Gly Leu Asp
 20 25 30

 Tyr Asp Tyr Thr Phe Asp Gly Asn Glu Glu Asp Lys Thr Glu Pro Ile
 35 40 45

 Asp Tyr Lys Asp Pro Cys Lys Ala Ala Val Phe Trp Gly Asp Ile Ala
 50 55 60

 Leu Asp Asp Glu Asp Leu Asn Ile Phe Gln Ile Asp Arg Thr Ile Asp
 65 70 75 80

 Leu Thr Gln Ser Pro Phe Gly Lys Leu Gly His Ile Thr Gly Gly Phe
 85 90 95

 Gly Asp His Gly Met Pro Lys Lys Arg Gly Ala Leu Tyr Gln Leu Ile
 100 105 110

 Glu Arg Ile Arg Arg Ile Gly Ser Gly Leu Glu Gln Asn Asn Thr Met
 115 120 125

 Lys Gly Lys Ala Pro Pro Lys Leu Ser Glu Gln Ser Glu Lys Asn Arg
 130 135 140

 Val Pro Arg Ala Ala Thr Ser Arg Thr Glu Arg Ile Trp Pro Gly Gly
 145 150 155 160

 Val Ile Pro Tyr Val Ile Gly Gly Asn Phe Thr Gly Ser Gln Arg Ala
 165 170 175

 Met Phe Lys Gln Ala Met Arg His Trp Glu Lys His Thr Cys Val Thr
 180 185 190

 Phe Thr Glu Arg Ser Asp Glu Glu Ser Tyr Ile Val Phe Thr Tyr Arg
 195 200 205

 Pro Cys Gly Cys Cys Ser Tyr Val Gly Arg Arg Gly Asn Gly Pro Gln
 210 215 220

 Ala Ile Ser Ile Gly Lys Asn Cys Asp Lys Phe Gly Ile Val Val His
 225 230 235 240

 Glu Leu Gly His Val Ile Gly Phe Trp His Glu His Thr Arg Pro Asp
 245 250 255

 Arg Asp Asn His Val Thr Ile Ile Arg Glu Asn Ile Gln Pro Gly Gln
 260 265 270

 Glu Tyr Asn Phe Leu Lys Met Glu Pro Gly Glu Val Asn Ser Leu Gly
 275 280 285

 Glu Arg Tyr Asp Phe Asp Ser Ile Met His Tyr Ala Arg Asn Thr Phe
 290 295 300

 Ser Arg Gly Met Phe Leu Asp Thr Ile Leu Pro Ser Arg Asp Asp Asn
 305 310 315 320

Gly Ile Arg Pro Ala Ile Gly Gln Arg Thr Arg Leu Ser Lys Gly Asp
 325 330 335
 Ile Ala Gln Ala Arg Lys Leu Tyr Arg Cys Pro Ala Cys Gly Glu Thr
 340 345 350
 Leu Gln Glu Ser Ser Gly Asn Leu Ser Ser Pro Gly Phe Pro Asn Gly
 355 360 365
 Tyr Pro Ser Tyr Thr His Cys Ile Trp Arg Val Ser Val Thr Pro Gly
 370 375 380
 Glu Lys Ile Val Leu Asn Phe Thr Thr Met Asp Leu Tyr Lys Ser Ser
 385 390 395 400
 Leu Cys Trp Tyr Asp Tyr Ile Glu Val Arg Asp Gly Tyr Trp Arg Lys
 405 410 415
 Ser Pro Leu Leu Gly Arg Phe Cys Gly Asp Lys Val Ala Gly Val Leu
 420 425 430
 Thr Ser Thr Asp Ser Arg Met Trp Ile Glu Phe Arg Ser Ser Ser Asn
 435 440 445
 Trp Val Gly Lys Gly Phe Ala Ala Val Tyr Glu Ala Ile Cys Gly Gly
 450 455 460
 Glu Ile Arg Lys Asn Glu Gly Gln Ile Gln Ser Pro Asn Tyr Pro Asp
 465 470 475 480
 Asp Tyr Arg Pro Met Lys Glu Cys Val Trp Lys Ile Met Val Ser Glu
 485 490 495
 Gly Tyr His Val Gly Leu Thr Phe Gln Ala Phe Glu Ile Glu Arg His
 500 505 510
 Asp Ser Cys Ala Tyr Asp His Leu Glu Val Arg Asp Gly Ala Ser Glu
 515 520 525
 Asn Ser Pro Leu Ile Gly Arg Phe Cys Gly Tyr Asp Lys Pro Glu Asp
 530 535 540
 Ile Arg Ser Thr Ser Asn Thr Leu Trp Met Lys Phe Val Ser Asp Gly
 545 550 555 560
 Thr Val Asn Lys Ala Gly Phe Ala Ala Asn Phe Phe Lys Glu Glu Asp
 565 570 575
 Glu Cys Ala Lys Pro Asp Arg Gly Gly Cys Glu Gln Arg Cys Leu Asn
 580 585 590
 Thr Leu Gly Ser Tyr Gln Cys Ala Cys Glu Pro Gly Tyr Glu Leu Gly
 595 600 605
 Pro Asp Arg Arg Ser Cys Glu Ala Ala Cys Gly Gly Leu Leu Thr Lys
 610 615 620

Leu Asn Gly Thr Ile Thr Thr Pro Gly Trp Pro Lys Glu Tyr Pro Pro
 625 630 635 640
 Asn Lys Asn Cys Val Trp Gln Val Ile Ala Pro Ser Gln Tyr Arg Ile
 645 650 655
 Ser Val Lys Phe Glu Phe Glu Leu Glu Gly Asn Glu Val Cys Lys
 660 665 670
 Tyr Asp Tyr Val Glu Ile Trp Ser Gly Pro Ser Ser Glu Ser Lys Leu
 675 680 685
 His Gly Lys Phe Cys Gly Ala Asp Ile Pro Glu Val Met Thr Ser His
 690 695 700
 Phe Asn Asn Met Arg Ile Glu Phe Lys Ser Asp Asn Thr Val Ser Lys
 705 710 715 720
 Lys Gly Phe Lys Ala His Phe Phe Ser Asp Lys Asp Glu Cys Ser Lys
 725 730 735
 Asp Asn Gly Gly Cys Gln His Glu Cys Val Asn Thr Met Gly Ser Tyr
 740 745 750
 Thr Cys Gln Cys Arg Asn Gly Phe Val Leu His Glu Asn Lys His Asp
 755 760 765
 Cys Lys Glu Ala Glu Cys Glu Gln Lys Ile His Ser Pro Ser Gly Leu
 770 775 780
 Ile Thr Ser Pro Asn Trp Pro Asp Lys Tyr Pro Ser Arg Lys Glu Cys
 785 790 795 800
 Thr Trp Val Ile Ser Ala Ile Pro Gly His Arg Ile Thr Leu Ala Phe
 805 810 815
 Asn Glu Phe Glu Val Glu Gln His Gln Glu Cys Ala Tyr Asp His Leu
 820 825 830
 Glu Ile Phe Asp Gly Glu Thr Glu Lys Ser Pro Ile Leu Gly Arg Leu
 835 840 845
 Cys Gly Ser Lys Ile Pro Asp Pro Leu Met Ala Thr Gly Asn Glu Met
 850 855 860
 Phe Ile Arg Phe Ile Ser Asp Ala Ser Val Gln Arg Lys Gly Phe Gln
 865 870 875 880
 Ala Thr His Ser Thr Glu Cys Gly Arg Leu Lys Ala Glu Ser Lys
 885 890 895
 Pro Arg Asp Leu Tyr Ser His Ala Gln Phe Gly Asp Asn Asn Tyr Pro
 900 905 910
 Gly Gln Leu Asp Cys Glu Trp Leu Leu Val Ser Glu Arg Gly Ser Arg
 915 920 925

Leu Glu Leu Ser Phe Gln Thr Phe Glu Val Glu Glu Ala Asp Cys
 930 935 940
 Gly Tyr Asp Tyr Val Glu Val Phe Asp Gly Leu Ser Ser Lys Ala Val
 945 950 955 960
 Gly Leu Gly Arg Phe Cys Gly Ser Gly Pro Pro Glu Glu Ile Tyr Ser
 965 970 975
 Ile Gly Asp Val Ala Leu Ile His Phe His Thr Asp Asp Thr Ile Asn
 980 985 990
 Lys Lys Gly Phe Tyr Ile Arg Tyr Lys Ser Ile Arg Tyr Pro Glu Thr
 995 1000 1005
 Met His Ala Lys Asn
 1010

<210> 96
 <211> 1012
 <212> PRT
 <213> Mus musculus

<400> 96
 Met Pro Leu Ala Thr Thr Leu Gly Thr Leu Val Leu Leu Leu Leu
 1 5 10 15
 Pro Leu Pro Arg Gly Ala Glu Val Thr Gly Asp His Ser Asn Val Ala
 20 25 30
 Leu Asp Tyr Gly Ala Leu Glu Gly Glu Glu Gly Thr Glu Gln Gln Leu
 35 40 45
 His Tyr His Asp Pro Cys Lys Ala Ala Val Phe Trp Gly Asp Ile Ala
 50 55 60
 Leu Asp Glu Asp Asp Leu Lys Leu Phe His Ile Asp Lys Ala Glu Asp
 65 70 75 80
 Trp Thr Lys Pro Ser Ile Asp Lys Pro Gly His Asp Thr Gly Gly Leu
 85 90 95
 Glu Glu Thr Ser Ala Arg Trp Pro Asn Asp Thr Ala Ser Asn Ala Ser
 100 105 110
 Ile Gln Ala Pro Arg Lys Asp Gly Lys Asp Ala Thr Thr Phe Leu Pro
 115 120 125
 Asn Pro Gly Thr Ser Asn Thr Thr Ala Lys Thr Phe Ser Ala Arg Val
 130 135 140
 Arg Arg Ala Thr Thr Ser Arg Thr Glu Arg Ile Trp Pro Gly Gly Val
 145 150 155 160
 Ile Pro Tyr Val Ile Gly Gly Asn Phe Thr Gly Thr Gln Arg Ala Ile

165	170	175
Phe Lys Gln Ala Met Arg His Trp Glu Lys His Thr Cys Val Thr Phe		
180	185	190
Val Glu Arg Thr Asp Glu Glu Ser Phe Ile Val Phe Ser Tyr Arg Thr		
195	200	205
Cys Gly Cys Cys Ser Tyr Val Gly Arg Arg Gly Gly Pro Gln Ala		
210	215	220
Ile Ser Ile Gly Lys Asn Cys Asp Lys Phe Gly Ile Val Ala His Glu		
225	230	235
Leu Gly His Val Val Gly Phe Trp His Glu His Thr Arg Pro Asp Arg		
245	250	255
Asp Gln His Val Thr Ile Ile Arg Glu Asn Ile Gln Pro Gly Gln Glu		
260	265	270
Tyr Asn Phe Leu Lys Met Glu Ala Gly Glu Val Ser Ser Leu Gly Glu		
275	280	285
Thr Tyr Asp Phe Asp Ser Ile Met His Tyr Ala Arg Asn Thr Phe Ser		
290	295	300
Arg Gly Val Phe Leu Asp Thr Ile Leu Pro Arg Arg Asp Asp Asn Gly		
305	310	315
Val Arg Pro Thr Ile Gly Gln Arg Val Arg Leu Ser Gln Gly Asp Ile		
325	330	335
Ala Gln Ala Arg Lys Leu Tyr Lys Cys Pro Ala Cys Gly Glu Thr Leu		
340	345	350
Gln Asp Thr Thr Gly Asn Phe Ser Ala Pro Gly Phe Pro Asn Gly Tyr		
355	360	365
Pro Ser Tyr Ser His Cys Val Trp Arg Ile Ser Val Thr Pro Gly Glu		
370	375	380
Lys Ile Ile Leu Asn Phe Thr Ser Met Asp Leu Phe Lys Ser Arg Leu		
385	390	395
400		
Cys Trp Tyr Asp Tyr Val Glu Ile Arg Asp Gly Tyr Trp Arg Lys Ala		
405	410	415
Pro Leu Leu Gly Arg Phe Cys Gly Asp Lys Ile Pro Glu Ser Leu Val		
420	425	430
Ser Ser Asp Ser Arg Leu Trp Val Glu Phe Arg Ser Ser Ser Ser		
435	440	445
Leu Gly Lys Gly Phe Phe Ala Val Tyr Glu Ala Met Cys Gly Gly Asp		
450	455	460
Ile Thr Lys Asp Ala Gly Gln Ile Gln Ser Pro Asn Tyr Pro Asp Asp		

465	470	475	480
Tyr Arg Pro Ser Lys Glu Cys Val Trp Arg Ile Thr Val Pro Asp Gly			
485	490	495	
Phe His Val Gly Leu Thr Phe Gln Ser Phe Glu Ile Glu Arg His Asp			
500	505	510	
Ser Cys Ala Tyr Asp Tyr Leu Glu Ile Arg Asp Gly Pro Thr Glu Asp			
515	520	525	
Ser Thr Leu Ile Gly His Phe Cys Gly Tyr Glu Lys Pro Glu Ala Val			
530	535	540	
Lys Ser Ser Ala Asn Arg Leu Trp Val Lys Phe Val Ser Asp Gly Ser			
545	550	555	560
Ile Asn Lys Ala Gly Phe Ala Ala Asn Phe Phe Lys Glu Val Asp Glu			
565	570	575	
Cys Ser Trp Pro Asp His Gly Gly Cys Glu Gln Arg Cys Val Asn Thr			
580	585	590	
Leu Gly Ser Tyr Thr Cys Ala Cys Asp Pro Gly Tyr Glu Leu Ala Ala			
595	600	605	
Asp Lys Lys Thr Cys Glu Val Ala Cys Gly Gly Phe Ile Thr Lys Leu			
610	615	620	
Asn Gly Thr Ile Thr Ser Pro Gly Trp Pro Lys Glu Tyr Pro Thr Asn			
625	630	635	640
Lys Asn Cys Val Trp Gln Val Val Ala Pro Val Gln Tyr Arg Ile Ser			
645	650	655	
Leu Gln Phe Glu Ala Phe Glu Leu Glu Gly Asn Asp Val Cys Lys Tyr			
660	665	670	
Asp Phe Val Glu Val Arg Ser Gly Leu Ser Pro Asp Ala Lys Leu His			
675	680	685	
Gly Lys Phe Cys Gly Ser Glu Thr Pro Glu Val Ile Thr Ser Gln Ser			
690	695	700	
Asn Asn Met Arg Val Glu Phe Lys Ser Asp Asn Thr Val Ser Lys Arg			
705	710	715	720
Gly Phe Arg Ala His Phe Phe Ser Asp Lys Asp Glu Cys Ala Lys Asp			
725	730	735	
Asn Gly Gly Cys Gln Gln Glu Cys Val Asn Thr Phe Gly Ser Tyr Leu			
740	745	750	
Cys Arg Cys Arg Asn Gly Tyr Arg Leu His Glu Asn Gly His Asp Cys			
755	760	765	
Lys Glu Ala Gly Cys Ala Tyr Lys Ile Ser Ser Ala Glu Gly Thr Leu			

770	775	780
Met Ser Pro Asn Trp Pro Asp Lys Tyr Pro Ser Arg Lys Glu Cys Thr		
785	790	795
Trp Asn Ile Ser Ser Thr Ala Gly His Arg Val Lys Ile Thr Phe Ser		
805	810	815
Glu Phe Glu Ile Glu Gln His Gln Glu Cys Ala Tyr Asp His Leu Glu		
820	825	830
Leu Tyr Asp Gly Thr Asp Ser Leu Ala Pro Ile Leu Gly Arg Phe Cys		
835	840	845
Gly Ser Lys Lys Pro Asp Pro Val Val Ala Thr Gly Ser Ser Leu Phe		
850	855	860
Leu Arg Phe Tyr Ser Asp Ala Ser Val Gln Arg Lys Gly Phe Gln Ala		
865	870	880
Val His Ser Thr Glu Cys Gly Arg Leu Lys Ala Glu Val Gln Thr		
885	890	895
Lys Glu Leu Tyr Ser His Ala Gln Phe Gly Asp Asn Asn Tyr Pro Ser		
900	905	910
Gln Ala Arg Cys Asp Trp Val Ile Val Ala Glu Asp Gly Tyr Gly Val		
915	920	925
Glu Leu Ile Phe Arg Thr Phe Glu Val Glu Glu Ala Asp Cys Gly		
930	935	940
Tyr Asp Phe Met Glu Ala Tyr Asp Gly Tyr Asp Ser Ser Ala Pro Arg		
945	950	960
Leu Gly Arg Phe Cys Gly Ser Gly Pro Leu Glu Glu Ile Tyr Ser Ala		
965	970	975
Gly Asp Ser Leu Met Ile Arg Phe His Thr Asp Asp Thr Ile Asn Lys		
980	985	990
Lys Gly Phe His Ala Arg Tyr Thr Ser Thr Lys Phe Gln Asp Ala Leu		
995	1000	1005
His Met Arg Lys		
1010		
<210> 97		
<211> 1015		
<212> PRT		
<213> Homo sapiens		
<400> 97		
Met Pro Arg Ala Thr Ala Leu Gly Ala Leu Val Ser Leu Leu Leu		
1	5	10
		15

Leu Pro Leu Pro Arg Gly Ala Gly Gly Leu Gly Glu Arg Pro Asp Ala
 20 25 30

 Thr Ala Asp Tyr Ser Glu Leu Asp Gly Glu Glu Gly Thr Glu Gln Gln
 35 40 45

 Leu Glu His Tyr His Asp Pro Cys Lys Ala Ala Val Phe Trp Gly Asp
 50 55 60

 Ile Ala Leu Asp Glu Asp Asp Leu Lys Leu Phe His Ile Asp Lys Ala
 65 70 80

 Arg Asp Trp Thr Lys Gln Thr Val Gly Ala Thr Gly His Ser Thr Gly
 85 90 95

 Gly Leu Glu Glu Gln Ala Ser Glu Ser Ser Pro Asp Thr Thr Ala Met
 100 105 110

 Asp Thr Gly Thr Lys Glu Ala Gly Lys Asp Gly Arg Glu Asn Thr Thr
 115 120 125

 Leu Leu His Ser Pro Gly Thr Leu His Ala Ala Ala Lys Thr Phe Ser
 130 135 140

 Pro Arg Val Arg Arg Ala Thr Thr Ser Arg Thr Glu Arg Ile Trp Pro
 145 150 155 160

 Gly Gly Val Ile Pro Tyr Val Ile Gly Gly Asn Phe Thr Gly Ser Gln
 165 170 175

 Arg Ala Ile Phe Lys Gln Ala Met Arg His Trp Glu Lys His Thr Cys
 180 185 190

 Val Thr Phe Ile Glu Arg Thr Asp Glu Glu Ser Phe Ile Val Phe Ser
 195 200 205

 Tyr Arg Thr Cys Gly Cys Cys Ser Tyr Val Gly Arg Arg Gly Gly Gly
 210 215 220

 Pro Gln Ala Ile Ser Ile Gly Lys Asn Cys Asp Lys Phe Gly Ile Val
 225 230 235 240

 Ala His Glu Leu Gly His Val Val Gly Phe Trp His Glu His Thr Arg
 245 250 255

 Pro Asp Arg Asp Gln His Val Thr Ile Ile Arg Glu Asn Ile Gln Pro
 260 265 270

 Gly Gln Glu Tyr Asn Phe Leu Lys Met Glu Ala Gly Glu Val Ser Ser
 275 280 285

 Leu Gly Glu Thr Tyr Asp Phe Asp Ser Ile Met His Tyr Ala Arg Asn
 290 295 300

 Thr Phe Ser Arg Gly Val Phe Leu Asp Thr Ile Leu Pro Arg Gln Asp
 305 310 315 320

Asp Asn Gly Val Arg Pro Thr Ile Gly Gln Arg Val Arg Leu Ser Gln
 325 330 335

 Gly Asp Ile Ala Gln Ala Arg Lys Leu Tyr Lys Cys Pro Ala Cys Gly
 340 345 350

 Glu Thr Leu Gln Asp Thr Thr Gly Asn Phe Ser Ala Pro Gly Phe Pro
 355 360 365

 Asn Gly Tyr Pro Ser Tyr Ser His Cys Val Trp Arg Ile Ser Val Thr
 370 375 380

 Pro Gly Glu Lys Ile Val Leu Asn Phe Thr Ser Met Asp Leu Phe Lys
 385 390 395 400

 Ser Arg Leu Cys Trp Tyr Asp Tyr Val Glu Val Arg Asp Gly Tyr Trp
 405 410 415

 Arg Lys Ala Pro Leu Leu Gly Arg Phe Cys Gly Asp Lys Ile Pro Glu
 420 425 430

 Pro Leu Val Ser Thr Asp Ser Arg Leu Trp Val Glu Phe Arg Ser Ser
 435 440 445

 Ser Asn Ile Leu Gly Lys Gly Phe Phe Ala Ala Tyr Glu Ala Thr Cys
 450 455 460

 Gly Gly Asp Met Asn Lys Asp Ala Gly Gln Ile Gln Ser Pro Asn Tyr
 465 470 475 480

 Pro Asp Asp Tyr Arg Pro Ser Lys Glu Cys Val Trp Arg Ile Thr Val
 485 490 495

 Ser Glu Gly Phe His Val Gly Leu Thr Phe Gln Ala Phe Glu Ile Glu
 500 505 510

 Arg His Asp Ser Cys Ala Tyr Asp Tyr Leu Glu Val Arg Asp Gly Pro
 515 520 525

 Thr Glu Glu Ser Ala Leu Ile Gly His Phe Cys Gly Tyr Glu Lys Pro
 530 535 540

 Glu Asp Val Lys Ser Ser Asn Arg Leu Trp Met Lys Phe Val Ser
 545 550 555 560

 Asp Gly Ser Ile Asn Lys Ala Gly Phe Ala Ala Asn Phe Phe Lys Glu
 565 570 575

 Val Asp Glu Cys Ser Trp Pro Asp His Gly Gly Cys Glu His Arg Cys
 580 585 590

 Val Asn Thr Leu Gly Ser Tyr Lys Cys Ala Cys Asp Pro Gly Tyr Glu
 595 600 605

 Leu Ala Ala Asp Lys Lys Met Cys Glu Val Ala Cys Gly Gly Phe Ile
 610 615 620

Thr Lys Leu Asn Gly Thr Ile Thr Ser Pro Gly Trp Pro Lys Glu Tyr
 625 630 635 640
 Pro Thr Asn Lys Asn Cys Val Trp Gln Val Val Ala Pro Ala Gln Tyr
 645 650 655
 Arg Ile Ser Leu Gln Phe Glu Val Phe Glu Leu Glu Gly Asn Asp Val
 660 665 670
 Cys Lys Tyr Asp Phe Val Glu Val Arg Ser Gly Leu Ser Pro Asp Ala
 675 680 685
 Lys Leu His Gly Arg Phe Cys Gly Ser Glu Thr Pro Glu Val Ile Thr
 690 695 700
 Ser Gln Ser Asn Asn Met Arg Val Glu Phe Lys Ser Asp Asn Thr Val
 705 710 715 720
 Ser Lys Arg Gly Phe Arg Ala His Phe Phe Ser Asp Lys Asp Glu Cys
 725 730 735
 Ala Lys Asp Asn Gly Gly Cys Gln His Glu Cys Val Asn Thr Phe Gly
 740 745 750
 Ser Tyr Leu Cys Arg Cys Arg Asn Gly Tyr Trp Leu His Glu Asn Gly
 755 760 765
 His Asp Cys Lys Glu Ala Gly Cys Ala His Lys Ile Ser Ser Val Glu
 770 775 780
 Gly Thr Leu Ala Ser Pro Asn Trp Pro Asp Lys Tyr Pro Ser Arg Arg
 785 790 795 800
 Glu Cys Thr Trp Asn Ile Ser Ser Thr Ala Gly His Arg Val Lys Leu
 805 810 815
 Thr Phe Asn Glu Phe Glu Ile Glu Gln His Gln Glu Cys Ala Tyr Asp
 820 825 830
 His Leu Glu Met Tyr Asp Gly Pro Asp Ser Leu Ala Pro Ile Leu Gly
 835 840 845
 Arg Phe Cys Gly Ser Lys Lys Pro Asp Pro Thr Val Ala Ser Gly Ser
 850 855 860
 Ser Met Phe Leu Arg Phe Tyr Ser Asp Ala Ser Val Gln Arg Lys Gly
 865 870 875 880
 Phe Gln Ala Val His Ser Thr Glu Cys Gly Gly Arg Leu Lys Ala Glu
 885 890 895
 Val Gln Thr Lys Glu Leu Tyr Ser His Ala Gln Phe Gly Asp Asn Asn
 900 905 910
 Tyr Pro Ser Glu Ala Arg Cys Asp Trp Val Ile Val Ala Glu Asp Gly
 915 920 925

Tyr Gly Val Glu Leu Thr Phe Arg Thr Phe Glu Val Glu Glu Ala
930 935 940

Asp Cys Gly Tyr Asp Tyr Met Glu Ala Tyr Asp Gly Tyr Asp Ser Ser
945 950 955 960

Ala Pro Arg Leu Gly Arg Phe Cys Gly Ser Gly Pro Leu Glu Ile
965 970 975

Tyr Ser Ala Gly Asp Ser Leu Met Ile Arg Phe Arg Thr Asp Asp Thr
980 985 990

Ile Asn Lys Lys Gly Phe His Ala Arg Tyr Thr Ser Thr Lys Phe Gln
995 1000 1005

Asp Ala Leu His Met Lys Lys
1010 1015

<210> 98

<211> 823

<212> PRT

<213> Homo sapiens

<400> 98

Met Pro Gly Val Ala Arg Leu Pro Leu Leu Leu Gly Leu Leu Leu
1 5 10 15

Pro Arg Pro Gly Arg Pro Leu Asp Leu Ala Asp Tyr Thr Tyr Asp Leu
20 25 30

Ala Glu Glu Asp Asp Ser Glu Pro Leu Asn Tyr Lys Asp Pro Cys Lys
35 40 45

Ala Ala Ala Phe Leu Gly Asp Ile Ala Leu Asp Glu Glu Asp Leu Arg
50 55 60

Ala Phe Gln Val Gln Gln Ala Val Asp Leu Arg Arg His Thr Ala Arg
65 70 75 80

Lys Ser Ser Ile Lys Ala Ala Val Pro Gly Asn Thr Ser Thr Pro Ser
85 90 95

Cys Gln Ser Thr Asn Gly Gln Pro Gln Arg Gly Ala Cys Gly Arg Trp
100 105 110

Arg Gly Arg Ser Arg Ser Arg Arg Ala Ala Thr Ser Arg Pro Glu Arg
115 120 125

Val Trp Pro Asp Gly Val Ile Pro Phe Val Ile Gly Gly Asn Phe Thr
130 135 140

Gly Ser Gln Arg Ala Val Phe Arg Gln Ala Met Arg His Trp Glu Lys
145 150 155 160

His Thr Cys Val Thr Phe Leu Glu Arg Thr Asp Glu Asp Ser Tyr Ile
165 170 175

Val Phe Thr Tyr Arg Pro Cys Gly Cys Cys Ser Tyr Val Gly Arg Arg
180 185 190

Gly Gly Gly Pro Gln Ala Ile Ser Ile Gly Lys Asn Cys Asp Lys Phe
195 200 205

Gly Ile Val Val His Glu Leu Gly His Val Val Gly Phe Trp His Glu
210 215 220

His Thr Arg Pro Asp Arg Asp Arg His Val Ser Ile Val Arg Glu Asn
225 230 235 240

Ile Gln Pro Gly Gln Glu Tyr Asn Phe Leu Lys Met Glu Pro Gln Glu
245 250 255

Val Glu Ser Leu Gly Glu Thr Tyr Asp Phe Asp Ser Ile Met His Tyr
260 265 270

Ala Arg Asn Thr Phe Ser Arg Gly Ile Phe Leu Asp Thr Ile Val Pro
275 280 285

Lys Tyr Glu Val Asn Gly Val Lys Pro Pro Ile Gly Gln Arg Thr Arg
290 295 300

Leu Ser Lys Gly Asp Ile Ala Gln Ala Arg Lys Leu Tyr Lys Cys Pro
305 310 315 320

Ala Cys Gly Glu Thr Leu Gln Asp Ser Thr Gly Asn Phe Ser Ser Pro
325 330 335

Glu Tyr Pro Asn Gly Tyr Ser Ala His Met His Cys Val Trp Arg Ile
340 345 350

Ser Val Thr Pro Gly Glu Lys Ile Ile Leu Asn Phe Thr Ser Leu Asp
355 360 365

Leu Tyr Arg Ser Arg Leu Cys Trp Tyr Asp Tyr Val Glu Val Arg Asp
370 375 380

Gly Phe Trp Arg Lys Ala Pro Leu Arg Gly Arg Phe Cys Gly Ser Lys
385 390 395 400

Leu Pro Glu Pro Ile Val Ser Thr Asp Ser Arg Leu Trp Val Glu Phe
405 410 415

Arg Ser Ser Ser Asn Trp Val Gly Lys Gly Phe Phe Ala Val Tyr Glu
420 425 430

Ala Ile Cys Gly Gly Asp Val Lys Lys Asp Tyr Gly His Ile Gln Ser
435 440 445

Pro Asn Tyr Pro Asp Asp Tyr Arg Pro Ser Lys Val Cys Ile Trp Arg
450 455 460

Ile Gln Val Ser Glu Gly Phe His Val Gly Leu Thr Phe Gln Ser Phe
465 470 475 480

Glu Ile Glu Arg His Asp Ser Cys Ala Tyr Asp Tyr Leu Glu Val Arg
485 490 495

Asp Gly His Ser Glu Ser Ser Thr Leu Ile Gly Arg Tyr Cys Gly Tyr
500 505 510

Glu Lys Pro Asp Asp Ile Lys Ser Thr Ser Ser Arg Leu Trp Leu Lys
515 520 525

Phe Val Ser Asp Gly Ser Ile Asn Lys Ala Gly Phe Ala Val Asn Phe
530 535 540

Phe Lys Glu Val Asp Glu Cys Ser Arg Pro Asn Arg Gly Gly Cys Glu
545 550 555 560

Gln Arg Cys Leu Asn Thr Leu Gly Ser Tyr Lys Cys Ser Cys Asp Pro
565 570 575

Gly Tyr Glu Leu Ala Pro Asp Lys Arg Arg Cys Glu Ala Ala Cys Gly
580 585 590

Gly Phe Leu Thr Lys Leu Asn Gly Ser Ile Thr Ser Pro Gly Trp Pro
595 600 605

Lys Glu Tyr Pro Pro Asn Lys Asn Cys Ile Trp Gln Leu Val Ala Pro
610 615 620

Thr Gln Tyr Arg Ile Ser Leu Gln Phe Asp Phe Phe Glu Thr Glu Gly
625 630 635 640

Asn Asp Val Cys Lys Tyr Asp Phe Val Glu Val Arg Ser Gly Leu Thr
645 650 655

Ala Asp Ser Lys Leu His Gly Lys Phe Cys Gly Ser Glu Lys Pro Glu
660 665 670

Val Ile Thr Ser Gln Tyr Asn Asn Met Arg Val Glu Phe Lys Ser Asp
675 680 685

Asn Thr Val Ser Lys Lys Gly Phe Lys Ala His Phe Phe Ser Val Leu
690 695 700

Glu Gly Ala Gly Asp Arg His Ser His Leu Ser Gly Leu Glu Leu Leu
705 710 715 720

Leu Cys Pro His Ala Leu Val Asp Thr Val Pro Ala Pro Pro Ser Ala
725 730 735

Leu His Gly Asp Thr His Ala His Thr His Thr His Val His Thr His
740 745 750

Cys Pro Ile Ala Gln Glu Thr Cys Arg Gly Pro Pro Leu Gly Ala Ser
755 760 765

Arg Leu Ser Pro Gln Gly Pro Gly His Leu Thr Leu Ala Pro Gln Glu
770 775 780

Gly Ser Tyr Leu Asp Phe Trp Asp Thr His Arg Gly Asp Pro Lys Pro
785 790 795 800

Arg Arg Arg Arg Lys Ser Leu Lys Thr Phe Ser Leu Thr Pro Ala Thr
805 810 815

Phe Arg Gly Ile Trp Ala Leu
820

<210> 99
<211> 1019
<212> PRT
<213> Xenopus laevis

<400> 99
Met Ser Cys Gly Ser Pro Gln Val Met Met Thr Leu Trp Thr Leu Thr
1 5 10 15

Cys Val Gly Leu Ile Leu Leu Gly Ala Ile Arg Leu Ser Leu Gly Leu
20 25 30

Asp Tyr Asp Leu Glu Ser Phe Asp Tyr Leu Met Glu Asp Asn Pro Glu
35 40 45

Glu Phe Asp Tyr Lys Asp Pro Cys Lys Ala Ala Ala Tyr Trp Gly Asp
50 55 60

Ile Ala Leu Asp Glu Asp Asp Leu Lys Trp Ile Phe Lys Asn Lys Ser
65 70 75 80

Asn Asp Leu Arg Asn Thr Arg His Asn Gln Thr His Pro Thr Thr Asp
85 90 95

Asn Phe Ser Glu Lys Leu Gly Thr Gly Ser Gln Asn Glu Thr Ser Ser
100 105 110

Asn Leu Asn Ser Lys Lys Val Lys Lys Gly Ser Arg Leu Lys Leu Leu
115 120 125

Ile Ala Glu Lys Ala Ala Thr Glu Thr Asn Ser Thr Phe Gln Val Gln
130 135 140

Thr Ser Asn Asp Arg Val Arg Arg Ala Ala Thr Ser Arg Thr Glu Arg
145 150 155 160

Ile Trp Pro Gly Gly Ile Ile Pro Tyr Ala Ile Ala Gly Asn Phe Thr
165 170 175

Gly Thr Gln Arg Ala Ile Phe Lys Gln Ala Met Arg His Trp Lys Lys
180 185 190

His Thr Cys Val Thr Phe Val Glu Arg Thr Asp Glu Glu Ser Phe Ile
195 200 205

Val Phe Thr Tyr Arg Pro Cys Gly Cys Cys Ser Tyr Val Gly Arg Arg

210	215	220													
Gly	Gly	Gly	Pro	Gln	Ala	Ile	Ser	Ile	Gly	Lys	Asn	Cys	Asp	Lys	Phe
225										235					240
Gly	Ile	Val	Val	His	Glu	Leu	Gly	His	Val	Val	Gly	Phe	Trp	His	Glu
	245								250					255	
His	Thr	Arg	Pro	Asp	Arg	Asp	Glu	His	Val	Ser	Ile	Ile	Arg	Glu	Asn
	260								265					270	
Ile	Gln	Pro	Gly	Gln	Glu	Tyr	Asn	Phe	Leu	Lys	Met	Glu	Pro	Gly	Glu
	275								280					285	
Val	Ser	Ser	Leu	Gly	Glu	Thr	Tyr	Asp	Phe	Asp	Ser	Ile	Met	His	Tyr
	290							295					300		
Ala	Arg	Asn	Thr	Phe	Ser	Arg	Gly	Val	Phe	Leu	Asp	Thr	Ile	Leu	Pro
	305										315				320
Arg	Arg	Ile	Asp	Thr	Ser	Val	Arg	Pro	Thr	Ile	Gly	Gln	Arg	Ile	Arg
				325					330					335	
Leu	Ser	Gln	Gly	Asp	Ile	Ala	Gln	Ala	Lys	Lys	Leu	Tyr	Lys	Cys	Pro
					340				345					350	
Ala	Cys	Gly	Glu	Thr	Leu	Gln	Asp	Ser	Ser	Gly	Asn	Phe	Ser	Ala	Pro
								355			360			365	
Gly	Tyr	Pro	Ser	Gly	Tyr	Pro	Ser	Tyr	Thr	His	Cys	Ile	Trp	Arg	Ile
								370			375			380	
Ser	Val	Thr	Pro	Gly	Glu	Lys	Ile	Ile	Leu	Asn	Phe	Thr	Thr	Met	Asp
										385			395		400
Leu	Phe	Lys	Ser	Arg	Leu	Cys	Trp	Tyr	Asp	Tyr	Ile	Glu	Ile	Arg	Asp
										405			410		415
Gly	Tyr	Trp	Arg	Lys	Ala	Ala	Leu	Leu	Gly	Arg	Leu	Cys	Gly	Asp	Lys
									420			425			430
Leu	Pro	Asp	Pro	Ile	Ile	Ser	Ser	Asp	Ser	Lys	Leu	Trp	Ile	Glu	Phe
									435				440		445
Arg	Ser	Ser	Ser	Asn	Ile	Leu	Gly	Lys	Gly	Phe	Phe	Ala	Ala	Tyr	Glu
									450			455			460
Ala	Ile	Cys	Gly	Gly	Asp	Ile	Lys	Lys	Asp	Ser	Gly	Gln	Ile	Gln	Ser
										465			470		480
Pro	Asn	Tyr	Pro	Asp	Asp	Tyr	Arg	Pro	Ala	Lys	Glu	Cys	Ile	Trp	Lys
										485			490		495
Ile	Thr	Val	Ser	Glu	Gly	Phe	Leu	Val	Gly	Leu	Ser	Phe	Gln	Ala	Phe
										500			505		510
Glu	Ile	Glu	Arg	His	Asp	Asn	Cys	Ala	Tyr	Asp	Tyr	Leu	Glu	Val	Arg

515	520	525
Asp Gly Phe Ser Glu Asp His Ala Leu Ile Gly Arg Phe Cys Gly Tyr		
530	535	540
Glu Lys Pro Glu Asp Ile Lys Ser Thr Ser Asn Lys Leu Trp Ile Lys		
545	550	555
Phe Ala Ser Asp Gly Ser Ile Asn Lys Ala Gly Phe Ser Ala Asn Phe		
565	570	575
Phe Lys Glu Met Asp Glu Cys Ser Arg Pro Asp Asn Gly Gly Cys Ser		
580	585	590
Gln Arg Cys Val Asn Thr Leu Gly Ser Tyr Lys Cys Val Cys Glu Pro		
595	600	605
Gly Phe Glu Leu Thr Ala Asp Lys Lys Ser Cys Glu Ala Ala Cys Gly		
610	615	620
Gly Phe Ile Thr Gln Leu Asn Gly Thr Ile Thr Ser Pro Gly Trp Pro		
625	630	635
Lys Glu Tyr Pro Thr Asn Lys Asn Cys Val Trp Gln Val Val Ala Pro		
645	650	655
Ala Gln Tyr Arg Ile Ser Leu Gln Phe Glu Val Phe Glu Leu Glu Gly		
660	665	670
Asn Asp Val Cys Lys Tyr Asp Tyr Leu Glu Ile Arg Ser Gly Leu Ser		
675	680	685
Ser Glu Ser Lys Leu His Gly Lys Phe Cys Gly Pro Glu Lys Pro Glu		
690	695	700
Val Ile Thr Ser Gln Gly Asn Thr Val Arg Ile Glu Phe Lys Ser Asp		
705	710	715
720		
Asn Thr Val Ser Lys Lys Gly Phe Lys Ala Asn Phe Phe Ser Asp Lys		
725	730	735
Asp Glu Cys Ser Lys Asp Asn Gly Gly Cys Gln His Asp Cys Val Asn		
740	745	750
Thr Phe Gly Ser Tyr Ile Cys Gln Cys Lys Asn Gly Phe Ile Leu His		
755	760	765
Glu Asn Gly His Asp Cys Lys Glu Ala Gly Cys Glu Gln Lys Leu Leu		
770	775	780
Asn Ala Glu Gly Thr Ile Ser Ser Pro Asn Trp Pro Glu Lys Tyr Pro		
785	790	795
Ser Arg Lys Glu Cys Thr Trp Asp Ile Ser Val Thr Ala Gly His Arg		
805	810	815
Val Lys Leu Val Phe Thr Asp Phe Glu Ile Glu Gln His Gln Glu Cys		

820	825	830
Ala Tyr Asp His Leu Glu Leu Tyr Asp Gly Pro Asn Gly Lys Ala Ala		
835	840	845
Ile Leu Gly Arg Phe Cys Gly Ser Lys Glu Pro Ser Pro Val Val Ala		
850	855	860
Ser Thr Asn Asn Met Phe Leu Arg Phe Tyr Ser Asp Ala Ser Val Gln		
865	870	875
Arg Lys Gly Phe Gln Ala Lys Tyr Ser Pro Glu Cys Gly Gly Arg Leu		
885	890	895
Lys Ala Glu Ile Gln Thr Asn Asp Ile Tyr Ser His Ala Gln Phe Gly		
900	905	910
Asp Asn Asn Tyr Pro Val Gln Ser Asn Cys Glu Trp Val Ile Val Ala		
915	920	925
Glu Asp Gly Tyr Gly Val Glu Leu Ile Phe Gln Thr Phe Glu Ile Glu		
930	935	940
Glu Glu Ser Asp Cys Gly Tyr Asp Tyr Met Glu Val Tyr Asp Gly Tyr		
945	950	955
Asp Ser Thr Ala Pro Arg Leu Gly Arg Tyr Cys Gly Ser Gly Pro Pro		
965	970	975
Glu Glu Met Tyr Ser Ala Gly Asp Ser Ile Met Ile Arg Phe His Thr		
980	985	990
Asp Asp Thr Ile Asn Lys Lys Gly Phe His Gly Gln Tyr Thr Ser Thr		
995	1000	1005
Lys Phe Gln Asp Ala Leu His Met Arg Arg Lys		
1010	1015	

<210> 100
 <211> 493
 <212> PRT
 <213> Rattus norvegicus

<400> 100			
Met Ala Asp Ser Lys Pro Leu Arg Thr Leu Asp Gly Asp Pro Val Ala			
1	5	10	15
Val Glu Ala Leu Leu Arg Asp Val Phe Gly Ile Val Val Asp Glu Ala			
20	25	30	
Ile Arg Lys Gly Thr Asn Ala Ser Glu Lys Val Cys Glu Trp Lys Glu			
35	40	45	
Pro Glu Glu Leu Lys Gln Leu Leu Asp Leu Glu Leu Gln Ser Gln Gly			
50	55	60	

Glu Ser Arg Glu Arg Ile Leu Glu Arg Cys Arg Ala Val Ile His Tyr
65 70 75 80

Ser Val Lys Thr Gly His Pro Arg Phe Phe Asn Gln Leu Phe Ser Gly
85 90 95

Leu Asp Pro His Ala Leu Ala Gly Arg Ile Ile Thr Glu Ser Leu Asn
100 105 110

Thr Ser Gln Tyr Thr Tyr Glu Ile Ala Pro Val Phe Val Leu Met Glu
115 120 125

Glu Glu Val Leu Lys Lys Leu Arg Ala Leu Val Gly Trp Asn Thr Gly
130 135 140

Asp Gly Val Phe Cys Pro Gly Gly Ser Ile Ser Asn Met Tyr Ala Ile
145 150 155 160

Asn Leu Ala Arg Phe Gln Arg Tyr Pro Asp Cys Lys Gln Arg Gly Leu
165 170 175

Arg Ala Leu Pro Pro Leu Ala Leu Phe Thr Ser Lys Glu Cys His Tyr
180 185 190

Ser Ile Thr Lys Gly Ala Ala Phe Leu Gly Leu Gly Thr Asp Ser Val
195 200 205

Arg Val Val Lys Ala Asp Glu Arg Gly Lys Met Ile Pro Glu Asp Leu
210 215 220

Glu Arg Gln Ile Ser Leu Ala Glu Ala Glu Gly Ser Val Pro Phe Leu
225 230 235 240

Val Ser Ala Thr Ser Gly Thr Thr Val Leu Gly Ala Phe Asp Pro Leu
245 250 255

Asp Ala Ile Ala Asp Val Cys Gln Arg His Gly Leu Trp Leu His Val
260 265 270

Asp Ala Ala Trp Gly Gly Ser Val Leu Leu Ser Arg Thr His Arg His
275 280 285

Leu Leu Asp Gly Ile Gln Arg Ala Asp Ser Val Ala Trp Asn Pro His
290 295 300

Lys Leu Leu Ala Ala Gly Leu Gln Cys Ser Ala Leu Leu Leu Arg Asp
305 310 315 320

Thr Ser Asn Leu Leu Lys Arg Cys His Gly Ser Gln Ala Ser Tyr Leu
325 330 335

Phe Gln Gln Asp Lys Phe Tyr Asn Val Ala Leu Asp Thr Gly Asp Lys
340 345 350

Val Val Gln Cys Gly Arg Arg Val Asp Cys Leu Lys Leu Trp Leu Met
355 360 365

Trp	Lys	Ala	Gln	Gly	Gly	Gln	Gly	Leu	Glu	Trp	Arg	Ile	Asp	Gln	Ala
370															
															380
Phe	Ala	Leu	Thr	Arg	Tyr	Leu	Val	Glu	Glu	Ile	Lys	Lys	Arg	Glu	Gly
385															
															400
Phe	Glu	Leu	Val	Met	Glu	Pro	Glu	Phe	Val	Asn	Val	Cys	Phe	Trp	Phe
405															
															415
Val	Pro	Pro	Ser	Leu	Arg	Gly	Lys	Lys	Glu	Ser	Pro	Asp	Tyr	Ser	Gln
420															
															430
Arg	Leu	Ser	Gln	Val	Ala	Pro	Val	Leu	Lys	Glu	Arg	Met	Val	Lys	Lys
435															
															445
Gly	Thr	Met	Met	Ile	Gly	Tyr	Gln	Pro	His	Gly	Thr	Arg	Ala	Asn	Phe
450															
															460
Phe	Arg	Met	Val	Val	Ala	Asn	Pro	Ile	Leu	Val	Gln	Ala	Asp	Ile	Asp
465															
															480
Phe	Leu	Leu	Gly	Glu	Leu	Glu	Arg	Leu	Gly	Gln	Asp	Leu			
485															

<210> 101

<211> 493

<212> PRT

<213> Mus musculus

<400> 101

Met	Ala	Asp	Ser	Lys	Pro	Leu	Arg	Thr	Leu	Asp	Gly	Asp	Pro	Val	Ala
1															
															15

Val	Glu	Ala	Leu	Leu	Gln	Asp	Val	Phe	Gly	Ile	Val	Val	Asp	Glu	Ala
20															
															30

Ile	Leu	Lys	Gly	Thr	Ser	Ala	Ser	Glu	Lys	Val	Cys	Glu	Trp	Lys	Glu
35															
															45

Pro	Glu	Glu	Leu	Lys	Gln	Leu	Leu	Asp	Leu	Glu	Leu	Gln	Ser	Gln	Gly
50															
															60

Glu	Ser	Arg	Glu	Gln	Ile	Leu	Glu	Arg	Cys	Arg	Thr	Val	Ile	His	Tyr
65															
															80

Ser	Val	Lys	Thr	Gly	His	Pro	Arg	Phe	Phe	Asn	Gln	Leu	Phe	Ser	Gly
85															
															95

Leu	Asp	Pro	His	Ala	Leu	Ala	Gly	Arg	Ile	Ile	Thr	Glu	Ser	Leu	Asn
100															
															110

Thr	Ser	Gln	Tyr	Thr	Tyr	Glu	Ile	Ala	Pro	Val	Phe	Val	Leu	Met	Glu
115															
															125

Glu	Glu	Val	Leu	Lys	Lys	Leu	Arg	Ala	Leu	Val	Gly	Trp	Asn	Ser	Gly
130															
															140

Asp Gly Val Phe Cys Pro Gly Gly Ser Ile Ser Asn Met Tyr Ala Met
 145 150 155 160
 Asn Leu Ala Arg Phe Gln Arg Tyr Pro Asp Cys Lys Gln Arg Gly Leu
 165 170 175
 Arg Ala Leu Pro Pro Leu Ala Leu Phe Thr Ser Lys Glu Cys His Tyr
 180 185 190
 Ser Ile Thr Lys Gly Ala Ala Phe Leu Gly Leu Gly Thr Asp Ser Val
 195 200 205
 Arg Val Val Lys Ala Asp Glu Arg Gly Arg Met Ile Pro Glu Asp Leu
 210 215 220
 Glu Arg Gln Ile Ile Leu Ala Glu Ala Glu Gly Ser Val Pro Phe Leu
 225 230 235 240
 Val Ser Ala Thr Ser Gly Thr Thr Val Leu Gly Ala Phe Asp Pro Leu
 245 250 255
 Asp Ala Ile Ala Asp Val Cys Gln Arg His Gly Leu Trp Phe His Val
 260 265 270
 Asp Ala Ala Trp Gly Gly Ser Val Leu Leu Ser Arg Thr His Arg His
 275 280 285
 Leu Leu Asp Gly Ile Gln Arg Ala Asp Ser Val Ala Trp Asn Pro His
 290 295 300
 Lys Leu Leu Ala Ala Gly Leu Gln Cys Ser Ala Leu Leu Leu Arg Asp
 305 310 315 320
 Thr Ser Asn Leu Leu Lys Arg Cys His Gly Ser Gln Ala Ser Tyr Leu
 325 330 335
 Phe Gln Gln Asp Lys Phe Tyr Asp Val Ala Leu Asp Thr Gly Asp Lys
 340 345 350
 Val Val Gln Cys Gly Arg Arg Val Asp Cys Leu Lys Leu Trp Leu Met
 355 360 365
 Trp Lys Ala Gln Gly Gly Gln Gly Leu Glu Arg Arg Ile Asp Gln Ala
 370 375 380
 Phe Ala Leu Thr Arg Tyr Leu Val Glu Glu Ile Lys Lys Arg Glu Gly
 385 390 395 400
 Phe Glu Leu Val Met Glu Pro Glu Phe Val Asn Val Cys Phe Trp Phe
 405 410 415
 Val Pro Pro Ser Leu Arg Gly Lys Lys Glu Ser Pro Asp Tyr Ser Gln
 420 425 430
 Arg Leu Ser Gln Val Ala Pro Val Leu Lys Glu Arg Met Val Lys Lys
 435 440 445

Gly Thr Met Met Ile Gly Tyr Gln Pro His Gly Thr Arg Ala Asn Phe
450 455 460

Phe Arg Met Val Val Ala Asn Pro Ile Leu Ala Gln Ala Asp Ile Asp
465 470 475 480

Phe Leu Leu Gly Glu Leu Glu Leu Leu Gly Gln Asp Leu
485 490

<210> 102

<211> 493

<212> PRT

<213> Homo sapiens

<400> 102

Met Ala Asp Ser Glu Ala Leu Pro Ser Leu Ala Gly Asp Pro Val Ala
1 5 10 15

Val Glu Ala Leu Leu Arg Ala Val Phe Gly Val Val Val Asp Glu Ala
20 25 30

Ile Gln Lys Gly Thr Ser Val Ser Gln Lys Val Cys Glu Trp Lys Glu
35 40 45

Pro Glu Glu Leu Lys Gln Leu Leu Asp Leu Glu Leu Arg Ser Gln Gly
50 55 60

Glu Ser Gln Lys Gln Ile Leu Glu Arg Cys Arg Ala Val Ile Arg Tyr
65 70 75 80

Ser Val Lys Thr Gly His Pro Arg Phe Phe Asn Gln Leu Phe Ser Gly
85 90 95

Leu Asp Pro His Ala Leu Ala Gly Arg Ile Ile Thr Glu Ser Leu Asn
100 105 110

Thr Ser Gln Tyr Thr Tyr Glu Ile Ala Pro Val Phe Val Leu Met Glu
115 120 125

Glu Glu Val Leu Arg Lys Leu Arg Ala Leu Val Gly Trp Ser Ser Gly
130 135 140

Asp Gly Ile Phe Cys Pro Gly Gly Ser Ile Ser Asn Met Tyr Ala Val
145 150 155 160

Asn Leu Ala Arg Tyr Gln Arg Tyr Pro Asp Cys Lys Gln Arg Gly Leu
165 170 175

Arg Thr Leu Pro Pro Leu Ala Leu Phe Thr Ser Lys Glu Cys His Tyr
180 185 190

Ser Ile Gln Lys Gly Ala Ala Phe Leu Gly Leu Gly Thr Asp Ser Val
195 200 205

Arg Val Val Lys Ala Asp Glu Arg Gly Lys Met Val Pro Glu Asp Leu

210	215	220
Glu Arg Gln Ile Gly Met Ala Glu Ala Glu Gly Ala Val Pro Phe Leu		
225	230	235
Val Ser Ala Thr Ser Gly Thr Thr Val Leu Gly Ala Phe Asp Pro Leu		
245	250	255
Glu Ala Ile Ala Asp Val Cys Gln Arg His Gly Leu Trp Leu His Val		
260	265	270
Asp Ala Ala Trp Gly Gly Ser Val Leu Leu Ser Gln Thr His Arg His		
275	280	285
Leu Leu Asp Gly Ile Gln Arg Ala Asp Ser Val Ala Trp Asn Pro His		
290	295	300
Lys Leu Leu Ala Ala Gly Leu Gln Cys Ser Ala Leu Leu Leu Gln Asp		
305	310	315
Thr Ser Asn Leu Leu Lys Arg Cys His Gly Ser Gln Ala Ser Tyr Leu		
325	330	335
Phe Gln Gln Asp Lys Phe Tyr Asp Val Ala Leu Asp Thr Gly Asp Lys		
340	345	350
Val Val Gln Cys Gly Arg Arg Val Asp Cys Leu Lys Leu Trp Leu Met		
355	360	365
Trp Lys Ala Gln Gly Asp Gln Gly Leu Glu Arg Arg Ile Asp Gln Ala		
370	375	380
Phe Val Leu Ala Arg Tyr Leu Val Glu Glu Met Lys Lys Arg Glu Gly		
385	390	395
400		
Phe Glu Leu Val Met Glu Pro Glu Phe Val Asn Val Cys Phe Trp Phe		
405	410	415
Val Pro Pro Ser Leu Arg Gly Lys Gln Glu Ser Pro Asp Tyr His Glu		
420	425	430
Arg Leu Ser Lys Val Ala Pro Val Leu Lys Glu Arg Met Val Lys Glu		
435	440	445
Gly Ser Met Met Ile Gly Tyr Gln Pro His Gly Thr Arg Gly Asn Phe		
450	455	460
Phe Arg Val Val Val Ala Asn Ser Ala Leu Thr Cys Ala Asp Met Asp		
465	470	475
480		
Phe Leu Leu Asn Glu Leu Glu Arg Leu Gly Gln Asp Leu		
485	490	

<210> 103
<211> 493
<212> PRT

<213> Homo sapiens

<400> 103

Met Ala Asp Ser Glu Ala Leu Pro Ser Leu Ala Gly Asp Pro Val Ala
1 5 10 15

Val Glu Ala Leu Leu Arg Ala Val Phe Gly Val Val Val Asp Glu Ala
20 25 30

Ile Gln Lys Gly Thr Ser Val Ser Gln Lys Val Cys Glu Trp Lys Glu
35 40 45

Pro Glu Glu Leu Lys Gln Leu Leu Asp Leu Glu Leu Arg Ser Gln Gly
50 55 60

Glu Ser Gln Lys Gln Ile Leu Glu Arg Cys Arg Ala Val Ile Arg Tyr
65 70 75 80

Ser Val Lys Thr Gly His Pro Arg Phe Phe Asn Gln Leu Phe Ser Gly
85 90 95

Leu Asp Pro His Ala Leu Ala Gly Arg Ile Ile Thr Glu Ser Leu Asn
100 105 110

Thr Ser Gln Tyr Thr Tyr Glu Ile Ala Pro Val Phe Val Leu Met Glu
115 120 125

Glu Glu Val Leu Arg Lys Leu Arg Ala Leu Val Gly Trp Ser Ser Gly
130 135 140

Asp Gly Ile Phe Cys Pro Gly Gly Ser Ile Ser Asn Met Tyr Ala Val
145 150 155 160

Asn Leu Ala Arg Tyr Gln Arg Tyr Pro Asp Cys Lys Gln Arg Gly Leu
165 170 175

Arg Thr Leu Pro Pro Leu Ala Leu Phe Thr Ser Lys Glu Cys His Tyr
180 185 190

Ser Ile Gln Lys Gly Ala Ala Phe Leu Gly Leu Gly Thr Asp Ser Val
195 200 205

Arg Val Val Lys Ala Asp Glu Arg Gly Lys Met Val Pro Glu Asp Leu
210 215 220

Glu Arg Gln Ile Gly Met Ala Glu Ala Glu Gly Ala Val Pro Phe Leu
225 230 235 240

Val Ser Ala Thr Ser Gly Thr Thr Val Leu Gly Ala Phe Asp Pro Leu
245 250 255

Gly Ala Ile Ala Asp Val Cys Gln Arg His Gly Leu Trp Leu His Val
260 265 270

Asp Ala Ala Trp Gly Gly Ser Val Leu Leu Ser Gln Thr His Arg His
275 280 285

Leu Leu Asp Gly Ile Gln Arg Ala Asp Ser Val Ala Trp Asn Pro His
 290 295 300

 Lys Leu Leu Ala Ala Gly Leu Gln Cys Ser Ala Leu Leu Gln Asp
 305 310 315 320

 Thr Ser Asn Leu Leu Lys Arg Cys His Gly Ser Gln Ala Ser Tyr Leu
 325 330 335

 Phe Gln Gln Asp Lys Phe Tyr Asp Val Ala Leu Asp Thr Gly Asp Lys
 340 345 350

 Val Val Gln Cys Gly Arg Arg Val Asp Cys Leu Lys Leu Trp Leu Met
 355 360 365

 Trp Lys Ala Gln Gly Asp Gln Gly Leu Glu Arg Arg Ile Asp Gln Ala
 370 375 380

 Phe Val Leu Ala Arg Tyr Leu Val Glu Glu Met Lys Lys Arg Glu Gly
 385 390 395 400

 Phe Glu Leu Val Met Glu Pro Glu Phe Val Asn Val Cys Phe Trp Phe
 405 410 415

 Val Pro Pro Ser Leu Arg Gly Lys Gln Glu Ser Pro Asp Tyr His Glu
 420 425 430

 Arg Leu Ser Lys Val Ala Pro Val Leu Lys Glu Arg Met Val Lys Glu
 435 440 445

 Gly Ser Met Met Ile Gly Tyr Gln Pro His Gly Thr Arg Gly Asn Phe
 450 455 460

 Phe Arg Val Val Ala Asn Ser Ala Leu Thr Cys Ala Asp Met Asp
 465 470 475 480

 Phe Leu Leu Asn Glu Leu Glu Arg Leu Gly Gln Asp Leu
 485 490

<210> 104
 <211> 493
 <212> PRT
 <213> Homo sapiens

<400> 104
 Met Ala Asp Ser Glu Ala Leu Pro Ser Leu Ala Gly Asp Pro Val Ala
 1 5 10 15

 Val Glu Ala Leu Leu Arg Ala Val Phe Gly Val Val Val Asp Glu Ala
 20 25 30

 Ile Gln Lys Gly Thr Ser Val Ser Gln Lys Val Cys Glu Trp Lys Glu
 35 40 45

 Pro Glu Glu Leu Lys Gln Leu Leu Asp Leu Glu Leu Arg Ser Gln Gly
 50 55 60

Glu Ser Gln Lys Gln Ile Leu Glu Arg Cys Arg Ala Val Ile Arg Tyr
 65 70 75 80
 Ser Val Lys Thr Gly His Pro Arg Phe Phe Asn Gln Leu Phe Ser Gly
 85 90 95
 Leu Asp Pro His Ala Leu Ala Gly Arg Ile Ile Thr Glu Ser Leu Asn
 100 105 110
 Thr Ser Gln Tyr Thr Tyr Glu Ile Ala Pro Val Phe Val Leu Met Glu
 115 120 125
 Glu Glu Val Leu Arg Lys Leu Arg Ala Leu Val Gly Trp Ser Ser Gly
 130 135 140
 Asp Gly Ile Phe Cys Pro Gly Gly Ser Ile Ser Asn Met Tyr Ala Val
 145 150 155 160
 Asn Leu Ala Arg Tyr Gln Arg Tyr Pro Asp Cys Lys Gln Arg Gly Leu
 165 170 175
 Arg Thr Leu Pro Pro Leu Ala Leu Phe Thr Ser Lys Glu Cys His Tyr
 180 185 190
 Ser Ile Gln Lys Gly Ala Ala Phe Leu Gly Leu Gly Thr Asp Ser Val
 195 200 205
 Arg Val Val Lys Ala Asp Glu Arg Gly Lys Met Val Pro Glu Asp Leu
 210 215 220
 Glu Arg Gln Ile Gly Met Ala Glu Ala Glu Gly Ala Val Pro Phe Leu
 225 230 235 240
 Val Ser Ala Thr Ser Gly Thr Thr Val Leu Gly Ala Phe Asp Pro Leu
 245 250 255
 Glu Ala Ile Ala Asp Val Cys Gln Arg His Gly Leu Trp Leu His Val
 260 265 270
 Asp Ala Ala Trp Gly Gly Ser Val Leu Leu Ser Gln Thr His Arg His
 275 280 285
 Leu Leu Asp Gly Ile Gln Arg Ala Asp Ser Val Ala Trp Asn Pro His
 290 295 300
 Lys Leu Leu Ala Ala Gly Leu Gln Cys Ser Ala Leu Leu Leu Gln Asp
 305 310 315 320
 Thr Ser Asn Leu Leu Lys Arg Cys His Gly Ser Gln Ala Ser Tyr Leu
 325 330 335
 Phe Gln Gln Asp Lys Phe Tyr Asp Val Ala Leu Asp Thr Gly Asp Lys
 340 345 350
 Val Val Gln Cys Gly Arg Arg Val Asp Cys Leu Lys Leu Trp Leu Met
 355 360 365

Trp Lys Ala Gln Gly Asp Gln Gly Pro Glu Arg Arg Ile Asp Gln Ala
370 375 380

Phe Val Leu Ala Arg Tyr Leu Val Glu Glu Met Lys Lys Arg Glu Gly
385 390 395 400

Phe Glu Leu Val Met Glu Pro Glu Phe Val Asn Val Cys Phe Trp Phe
405 410 415

Val Pro Pro Ser Leu Arg Gly Lys Gln Glu Ser Pro Asp Tyr His Glu
420 425 430

Arg Leu Ser Lys Val Ala Pro Val Leu Lys Glu Arg Met Val Lys Glu
435 440 445

Gly Ser Met Met Ile Gly Tyr Gln Pro His Gly Thr Arg Gly Asn Phe
450 455 460

Phe Arg Val Val Val Ala Asn Ser Ala Leu Thr Cys Ala Asp Met Asp
465 470 475 480

Phe Leu Leu Asn Glu Leu Glu Arg Leu Gly Gln Asp Leu
485 490

<210> 105

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:oligonucleotide primer

<400> 105

ccatgtggca gctgaggc tt cat

23

<210> 106

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:oligonucleotide primer

<400> 106

aaagccccag gtcctttgc tagct

25

<210> 107

<211> 27

<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 107
ggatgaacca gactttgaat agcagtg

27

<210> 108
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 108
ggctctcaag ccccccatttc

20

<210> 109
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 109
atgcgaagtc actcttacct ctgatgat

28

<210> 110
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 110
gggagctgat cttgagttat ttaacatagc

30

<210> 111
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 111

ctgaatggaa ccatcaccag c	21
<210> 112	
<211> 26	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide primer	
<400> 112	
atcagcacta tttcttcatg tgcagg	26
<210> 113	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide primer	
<400> 113	
ctgcacttgg ctggaactta	20
<210> 114	
<211> 26	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide primer	
<400> 114	
tttcatctcc tactccaggt gtacca	26
<210> 115	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide primer	
<400> 115	
atctccacag gccctgtaat	20
<210> 116	
<211> 21	

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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
      primer

<400> 116
ccagccatgc tcagagtgac t                                21

<210> 117
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
      primer

<400> 117
ttgccaacag caaggtcaat gccac                                25

<210> 118
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
      primer

<400> 118
cgccactgtg gtcgatcat                                19

<210> 119
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
      primer

<400> 119
ccacagtttg ggatacagaa caatt                                25

<210> 120
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide

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primer

<400> 120
actgtgcttc caacacctggta gccctga 27

<210> 121
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide primer

<400> 121
agcctttttg acaaacggaag ag 22

<210> 122
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide primer

<400> 122
aagctggagt atcaggccta tga 23

<210> 123
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide primer

<400> 123
agagtgccca gaccaggcgc ccttt 25

<210> 124
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide primer

<400> 124
ctccccctcac aatgccattg 20

<210> 125	
<211> 22	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 125	
ctgaagctgg catggtaactt aa	22
<210> 126	
<211> 26	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 126	
cagtcctttc tctttgaacg acagcg	26
<210> 127	
<211> 22	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 127	
ttgttaggctt caatctttc ca	22
<210> 128	
<211> 21	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 128	
tgctatgaaa ggcaaggata a	21
<210> 129	
<211> 29	
<212> DNA	
<213> Artificial Sequence	

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 129
tgaatgccac aactttatca aagtatttg

29

<210> 130
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 130
aaaaccatct catcgttct tg

22

<210> 131
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 131
cagaagctgt tccgagaagt c

21

<210> 132
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 132
atgaagggcc taaaccaccc caacat

26

<210> 133
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 133	
caatcacctc aaagagcttc ac	22
<210> 134	
<211> 19	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 134	
agaacacccccc tggctcccta	19
<210> 135	
<211> 24	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 135	
acaccaggcct gtgaccctgg ctat	24
<210> 136	
<211> 22	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 136	
gtttcacact cgttcacatc ct	22
<210> 137	
<211> 19	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 137	
gcaacagcat ggtgatctg	19
<210> 138	

<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide primer

<400> 138
cttcgaatg cacaggaacc ccttct 26

<210> 139
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide primer

<400> 139
cgccaggttg aggatataga t 21

<210> 140
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide primer

<400> 140
ctgcaaccac atgatcatac aa 22

<210> 141
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide primer

<400> 141
atcaggaaac ctgaccacac ttgtaa 26

<210> 142
<211> 22
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 142
atggatgaag acatgctcct tt

22

<210> 143
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 143
caattttggc tctggagaaa ga

22

<210> 144
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 144
tctcagtgcc gatggacctc atagaa

26

<210> 145
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 145
cagtgaagca cataggtggc tt

22

<210> 146
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:oligonucleotide
primer

<400> 146
ctgggagaga catacgactt tg

22

<210> 147	
<211> 25	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 147	
cccggaacac cttctcaaga ggagt	25
<210> 148	
<211> 22	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 148	
gggaaggatg gtgtctaaga aa	22
<210> 149	
<211> 19	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 149	
tgtgctttg ttagccaga	19
<210> 150	
<211> 29	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:oligonucleotide	
primer	
<400> 150	
catcaatctg cttgctacac ttctcacca	29
<210> 151	
<211> 16	
<212> DNA	

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:oligonucleotide primer

<400> 151
ccaaagccct cggaac 16

<210> 152
<211> 22
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:oligonucleotide primer

<400> 152
tgtcgggatg tcatacacta ca 22

<210> 153
<211> 26
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:oligonucleotide primer

<400> 153
tgtcaaaaaca gaccacccaa gatttt 26

<210> 154
<211> 22
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:oligonucleotide primer

<400> 154
atcaagtcca gcataacaatt gg 22

<210> 155
<211> 38
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
oligonucleotide primer

<400> 155	
aagctttgtc agcgcaaccc ccaggtctgc ggcccagg	38
<210> 156	
<211> 37	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:	
oligonucleotide primer	
<400> 156	
ctcgagacag cgtccagtca tgggtcaaaa ctcttcc	37
<210> 157	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:	
oligonucleotide primer	
<400> 157	
gagaacacgc caggcagtt	20
<210> 158	
<211> 21	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:	
oligonucleotide primer	
<400> 158	
ctcccttcac tgtgcctgcc c	21
<210> 159	
<211> 19	
<212> DNA	
<213> Artificial Sequence	
<220>	
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